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# 64 HEMATOPOIETIC STEM CELL MULTIPLIER.

(57) A hematopoietic stem cell multiplier containing at least one hepatocyte proliferating factor as an active ingredient. Because hepatocyte proliferating factors have an activity of proliferating undifferentiated pluripotent hematopoietic stem cells, they are useful as a hematopoietic stem cell multiplier not only in treating bone marrow inhibition and insufficiency but also in effecting in vitro growth of peripheral blood stem cells and bone marrow stem cells. Further it is possible to derive a hepatocyte proliferating factor from human normal fibroblasts among other hepatocyte proliferating factors by the gene recombination technique, and the obtained proliferating factor is, too, useful as a hematopoietic stem cell multiplier.

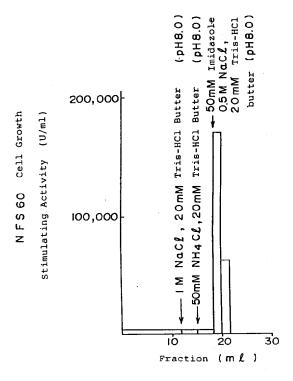


Figure 6

#### FIELD OF INVENTION

The present invention relates to a hemopoietic stem cell augmenting agent. In particular, the present invention relates to a hemopoietic stem cell augmenting agent which has an activity to augment undifferentiated pluripotent hemopoietic stem cells. According to the present invention, therapeutic agents to restore bone marrow growth after the use of anti-cancer agents or after bone marrow transplantation or therapeutic agents for bone marrow malfunctions such as hypoplastic anemia and osteomyelodysplasia syndrome can be provided. Furthermore, this hemopoietic factor having the hemopoietic stem cell augmenting activity can be used as a useful reagent or the like for in vitro proliferation of peripheral stem cells and bone marrow stem cells and furthermore as an analytical reagent or as an antigen for preparation of antibodies.

Furthermore, the present invention also relates to a novel protein which belongs to hepatocyte growth factors and has a hemopoietic stem cell augmenting activity.

#### BACKGROUND OF THE INVENTION

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Recently, it has been found that in the differentiation of undifferentiated pluripotent hemopoietic stem cells to mature blood cells, a number of hemopoietic factors are mutually involved in various levels and thus a complicated hemopoietic network system is formed. Furthermore, most of the hemopoietic factors are being genetically cloned and several hemopoietic factors are being mass produced by genetic recombination techniques, and their clinical applications are in progress. On the other hand, although pluripotent hemopoietic stem cells are characterized by their self-replicating ability (growth), sufficient studies have not been done on the growth factors which affect the undifferentiated pluripotent hemopoietic stem cells in the bone marrow.

It is known that bone marrow stromal cells play a major role in the growth of the pluripotent hemopoietic stem cells and their differentiation into mature cells; some kind of fluid factors secreted by the stromal cells, or intercellular reactions or the like are thought to be involved in hemopoieses in the bone marrow.

For example, it is known that bone marrow stromal cell line MC3T3-G2/PA-6(PA-6) which is established from the calvaria of C57B1/6 newborn mouse supports the growth of mouse pluripotent hemopoietic stem cells (Kodama, H. et al., J. Cell. Physiol., 112, 89, 1982).

In recent years, a ligand for a c-kit protein, which is a tyrosine kinase receptor being expressed in pluripotent stem cells, has attracted attention as a factor involved in the growth of undifferentiated stem cells and great efforts have been made to substantiate the ligand; in 1990, three research groups succeeded in genetically cloning the ligand. They were reported as SCF (stem cell factor; K. M. Zsebo et al., Cell, 63, 195-201, 1990), MGF (mast cell growth factor; D. E. Williams et al., Cell, 63, 167-174, 1990) and KL (c-kit ligand; Huang et al., Cell, 63, 225-233, 1990).

At present, the mechanism of the action of the ligand is being studied using the c-kit ligand which is mass produced by genetic recombination techniques and studies to date are clarifying that this factor acts on the stem cells which are differentiated to a certain extent (Hayashi et al., Int. J. Hematology, Suppl. No. 1, p198, 1991).

Accordingly, it is now considered that aside from this protein, other factors exist which act on pluripotent hemopoietic stem cells which are differentiated to a lesser extent in the bone marrow.

The hemopoietic factors having the above-mentioned activities can be made into useful pharmaceuticals such as therapeutic agents to restore bone marrow growth after the use of anti-cancer agents or after bone marrow transplantation and therapeutic agents for bone marrow malfunctions such as hypoplastic anemia and osteomyelodysplasia syndrome.

Furthermore, the hemopoietic factors having the above-mentioned activities can be used as a useful reagent or the like for in vitro proliferation of peripheral stem cells and bone marrow stem cells and furthermore as an analytical reagent or as an antigen for preparation of antibodies.

#### DISCLOSURE OF THE INVENTION

It is an object of the present invention to provide a hemopoietic stem cell augmenting agent which contains at least one hepatocyte growth factor as an active component. In particular, an object of the present invention is to provide a hemopoietic stem cell augmenting agent which has an activity for proliferating undifferentiated pluripotent hemopoietic stem cells. This hemopoietic stem cell augmenting agent eventually promotes proliferation of osteoclasts, the progeny of hemopoietic stem cells, as well as various blood cells. This hemopoietic stem cell augmenting agent may contain interleukin 3 and/or interleukin 7 as an active component(s) in addition to at least one of the hepatocyte growth factors.

A further object of the present invention is to provide a novel protein which is a type of hepatocyte growth factor and has a hemopoietic stem cell augmenting activity. In particular, such protein can be obtained from a culture fluid of human normal fibroblasts or by genetic recombination techniques using human normal fibroblasts as a gene supply source. It is a particular object of the present invention to provide a recombinant human hepatocyte growth factor protein, which is one type of hepatocyte growth factor and has the amino acid sequence of SEQ ID NO: 2 shown in Sequence Listing or a substance which has equivalent effectiveness. The protein is derived from human normal fibroblasts and preferably from fibroblasts of the highest normality. The present invention also provides a DNA which is characterized in that it contains a nucleotide sequence encoding an amino acid sequence of a protein, which is a type of hepatocyte growth factor and has a hemopoietic stem cell augmenting activity, or a substance equivalent in terms of effectiveness; a recombinant expression vector in which said coding nucleotide sequence is so incorporated as to be expressed; a transformant which is obtained by the transformation of a host cell using the expression vector; and a process for the preparation of a recombinant human hepatocyte growth factor protein, which is one of hepatocyte growth factors and has a hemopoietic stem cell augmenting activity, which is characterized in that the transformant is cultured under the conditions in which said protein can be expressed in a culture medium and the recombinant human hepatocyte growth factor protein, which is one type of hepatocyte growth factor and has a hemopoietic stem cell augmenting activity, is prepared from the culture. In particular, the present invention relates to a recombinant which is derived from human normal fibroblasts and considered preferably from fibroblasts of the highest normality, and a process for the preparation of the same and use of the same.

An agent which contains at least one type of hepatocyte growth factor disclosed in the present invention as an active component is effective for use as a therapeutic agent for bone marrow suppression or as a therapeutic agent for bone marrow malfunctions.

An agent which contains at least one type of hepatocyte growth factor disclosed in the present invention as an active component is effective for in vitro growth of peripheral blood stem cells and bone marrow stem cells, and thus the present invention also provides a process for in vitro proliferation or the cultivation of stem cells using said agent.

#### BRIEF DESCRIPTION OF THE DRAWINGS

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Figure 1 illustrates a construction of an expression vector pSRalphaBX.

Figure 2 is a nucleotide sequence and an amino acid sequence of HGF cDNA derived from human normal fibroblasts.

Figure 3 illustrates a construction of a human HGF expression vector pSRalphaFDF-1 for the expression of mammalian cells.

Figure 4 illustrates the relationships between HGF activity and fractions of a DEAE Sephacel eluent obtained from the culture supernatant of human normal fibroblast hepatocyte growth factor producing monkey COS-1 cells which are obtained in Example 3(6).

Figure 5 illustrates the relationships between HGF activity and fractions of a heparin eluent of heparin-Sepharose chromatography of a culture supernatant of human normal fibroblast hepatocyte growth factor producing monkey COS-1 cells which are obtained in Example 3(6).

Figure 6 illustrates the relationships between HGF activity and fractions of a zinc eluent of zinc chelate affinity chromatography of the culture supernatant of human normal fibroblast hepatocyte growth factor producing monkey COS-1 cells which are obtained in Example 3(6).

Figure 7 is a SDS-polyacrylamide gel electrophoresis pattern of recombinant-type human HGF obtained by the purification process as described in Example 4, under non-reducing conditions.

Figure 8 illustrates NFS60 cell proliferating activity of HGF derived from human placenta and purified HGF derived from fibroblasts.

Figure 9 is a chromatogram of reversed-phase high performance liquid chromatography as described in Example 1.

#### BEST MODE FOR CARRYING OUT THE INVENTION

The present inventors focused their attentions on the fact that proliferation of cells of immature myeloblast cell line NFS 60 (Kevin L. Holmes et al., Proc. Natl. Acad. Sci. USA, 82, 6687-6691, 1985) is dependent on interleukin 3 (IL-3) and that among the hemopoietic factors so far known, IL-3 is one of the cytokinins acting on undifferentiated hemopoietic stem cells and intensively searched for hemopoietic stem cell augmenting factors other than IL-3, using the augmenting activity on NFS60. As a result, the present

inventors discovered a factor, which has hemopoietic stem cell augmenting activity, in the fibroblasts which are known to produce hemopoietic factors such as interleukin 1 (IL-1), interleukin 6 (IL-6), interleukin 7 (IL-7), interleukin 8 (IL-8), interleukin 11 (IL-11) and the c-kit ligand and found that the factor is a novel factor which is a type of hepatocyte growth factor. Subsequently, the present inventors also found that this growth factor supports the growth of hemopoietic stem cells in an evaluation system using human bone marrow cells and murine bone marrow cells.

This novel factor is a novel physiologically active protein which has an activity to support the growth of pluripotent hemopoietic stem cells and has a molecular weight of about 60,000 and an N-terminal amino acid sequence of SEQ ID NO: 1 shown in Sequence Listing.

Furthermore, this novel factor has an amino acid composition shown in Table 1. Since the N-terminal amino acid sequence and the amino acid composition of the physiologically active protein of this novel factor are different from those of known proteins, this belongs to a novel hemapoietic factor.

The novel factor of the present invention can be obtained in a purified form as a physiologically active protein using a culture fluid obtained by culturing human normal fibroblasts as the starting material and using a combination of various types of chromatography.

Details of physico-chemical properties and biological properties of the physiologically active protein of the present invention thus obtained are as follows:

(1) Molecular weight:

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- 60,000 (by SDS-polyacrylamide gel electrophoresis; Laemmli U. K., Nature, 227, 680-685, 1970)
- (2) N-terminal amino acid sequence (16 residues).
  - Shown in SEQ ID NO: 1 in Sequence Listing.
- (3) Amino acid composition
  - Shown in Table 1.
- (4) Biological activities
  - A. Exhibits an augmenting activity on pluripotent myeloblasts derived from mice (NFS60).
  - B. Exhibits an augmenting activity in combination with IL-3 or with IL-3 and interleukin-7 (IL-7) on bone marrow cells of 5-fluorouracil (5-FU)-treated mice.
  - C. Exhibits an augmenting activity on hemopoietic stem cells derived from human normal bone marrow.

On the other hand, since this factor having the hemopoietic stem cell augmenting activity is a type of hepatocyte growth factor, the present inventors studied a growth factor which was reported as one type of hepatocyte growth factor and a typical hepatocyte growth factor (Nakamura, T. et al., Biochem. Biophys. Res. Commun., 122, 1450-1459, 1984) and considered this factor to have an proliferating activity on NFS60 strain. Furthermore, they also considered this augmenting factor to support the growth of hemopoietic stem cells in an evaluation system using human and murine bone marrow cells or to have the hemopoietic stem cell augmenting activity.

Further, some of the hepatocyte growth factors are reported to have physiological activities other than the augmenting activity on hepatic parenchyma cells, such as the mitogenic activity on epithelial cells (Gherardi, E. et al., Nature, 346, 228, 1990) and a cytotoxic activity on tumor cells (Higashio, K. et al., Biochem. Biophys. Res. Commun., 170, 397-404, 1990); however, an action on bone marrow hemopoietic stem cells was not known to date.

Furthermore, since a typical hepatocyte growth factor derived from the liver, one of the hepatocyte growth factors, was genetically cloned and the whole nucleotide sequence was determined (Nakamura et al., Nature, 342, 440-443, 1989), that may used as a hemopoietic stem cell augmenting agent in the present invention. Furthermore, the factor, a type of hepatocyte growth factor and used in the present invention as a hemopoietic stem cell augmenting agent, can also be produced by culturing cells such as human normal fibroblasts. Furthermore, the factor can be produced by taking out a gene from human normal fibroblasts or the like, applying genetic recombination technique to the gene thus obtained and culturing the resultant transformants.

Thus, according to the present invention, the novel factor can be obtained also using genetic recombination technique. This novel factor is characterized in that it has the amino acid sequence of SEQ ID NO: 2 in Sequence Listing. This novel factor can be obtained using a cloned cDNA as specifically explained hereinbelow.

The above-mentioned factor can be obtained in a purified form, for example, from a culture fluid obtained by culturing human normal fibroblasts as the starting material by means of using a combination of various kinds of chromatography.

Cells such as human normal fibroblasts can be grown or cultured using various kinds of conventional cell culture media. Examples of such media are those in which substances such as a carbon source, a

nitrogen source, vitamins, amino acids and nucleotides are contained and substances selected from the group consisting of meat broth, peptone, casamino acid, yeast extract, fish meal extract, potatoes, malt juice, cow's milk, blood, serum, hormones, antibiotics and cell growth factors are added if desired; media which are widely used in general and are commercially available can be used as they are or with appropriate modifications.

Examples of these media include RPMI-1640 medium, MEM medium, Dulbecco-Eagle medium, DMEM medium, McCoy 5A medium, Iscove's modified Dulbecco's medium and Ham F12 medium.

In the proliferation and cultivation of cells such as human normal fibroblasts as mentioned above, optimal conditions for the growth of said human normal fibroblasts, such as pH, temperature, aeration, stirring and frequency of medium exchange, can be appropriately determined, for example, by experiments.

In the cultivation of human normal fibroblasts, cell adhesion factors such as collagen, fibronectin, gelatin, poly-L-lysine and poly-D-lysine can be added or microcarrier beads made of dextran, polyacrylamide, polystyrene, gelatin, glass or the like can be used, if appropriate.

If desired, the human normal fibroblasts obtained by the above-mentioned cell proliferation or cultivation are subjected to induction, for example, by a treatment with conventional inducing agents such as poly I/C to induce the production of the novel protein which is a type of hepatocyte growth factor and has a hemopoietic stem cell augmenting activity; the protein of the present invention thus obtained is isolated by a conventional procedure.

Examples of isolation procedures include a method by ultrasonic destruction, mechanical destruction or freezing and thawing, a method by osmotic shock or the like and an isolation method in which a culture supernatant is precipitated, for example, using a protein precipitant.

In addition to the above-mentioned separation methods, the targeted protein can be purified by various kinds of isolation and purification methods which are widely used in general, utilizing physical properties and chemical properties of the protein.

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Examples of these isolation and purification methods include solubilization by homogenization or sonication, extractions with buffer solutions containing various kinds of salts, solubilization or precipitation with acids or alkalis, or else extraction or precipitation with organic solvents, salting out by precipitation with protein precipitants such as ammonium sulfate, ultrafiltrations by dialysis or ultra-filtration using a membrane filter, adsorption chromatography, gel filtration chromatography or the like, ion exchange chromatography, reversed-phase chromatography, affinity chromatography, counter-current distribution chromatography, high-performance liquid chromatography, isoelectric point or gel electrophoresis; these methods are used alone or in combination if appropriate.

If the protein is derived from human normal fibroblasts, affinity chromatography using a carrier bound with a blue pigment (a blue gel carrier), a carrier bound with zinc via chelate bonds (a zinc chelate gel carrier), a carrier bound with heparin (a heparin gel carrier), a carrier bound with antibodies or the like can be advantageously used. In particular, the zinc chelate gel carrier (media), the blue gel carrier (media) and the heparin gel carrier (media) are preferably used. A method in which the zinc chelate gel carrier and the heparin gel carrier are used in combination is more preferably used; the order of the use of the carriers is not particularly restricted but the use of the zinc chelate gel carrier after the heparin gel carrier is particularly preferable.

An example of the zinc chelate gel carrier used in the present invention is a carrier in which exchangeable groups having chelating capacity, such as a bis-carboxymethylimino group [N(CH<sub>2</sub>COOH)<sub>2</sub>], are bound to agarose, cellulose, polyacrylamide gel or the like is treated in a solution of zinc salt such as zinc chloride. Preferably, a carrier in which zinc is bound via chelate bonds to an insoluble polysaccharide gel carrier, such as "Chelating Sepharose" (a product of Pharmacia) is used.

Purification of hepatocyte growth factors using the zinc chelate gel carrier is carried out as follows. Namely, first, a solution containing the hepatocyte growth factors is made to come into contact with and adsorbed onto the zinc chelate gel carrier. The adsorption can be done either by a batch method or a column method but the column method gives higher adsorption efficiency.

The elution can be carried out using an acidic buffer solution such as phosphoric acid, acetic acid and citric acid and preferably at pH 5 or below. However, in high ionic strength, elution at higher pHs is possible. Furthermore, a gradient elution in which a concentration of imidazole, histamine, glycine or ammonium chloride is gradually increased gives good results. It is also possible to use a method in which metal ions are depleted from gels using a chelating agent such as EGTA or EDTA.

The ionic strength can be increased by increasing the concentration of acids such as phosphoric acid, acetic acid and citric acid in a buffer solution or by adding neutral salts such as sodium chloride and potassium chloride (0.2 to 1.0 M).

The composition and volume of the eluent are not particularly restricted and the optimal conditions for elution are appropriately determined depending on co-existing proteins or contaminants, the amount of hepatocyte growth factors, a column size or the like.

The heparin gel carrier used in the present invention can be any insoluble carrier formed from a synthetic polymer, a polysaccharide in which cellulose, agarose or the like is the starting material and forms the backbone molecule, or the like to which heparin is bound. Examples of the heparin gel carrier include "Heparin-Sepharose CL-6B (a product of Pharmacia), "Heparin Toyopearl" (a product of Toso) and "Heparin Cellulofine (a product of Chisso).

In the case where a solution containing hepatocyte growth factors is allowed to come into contact with the heparin gel carrier, it is desirable to control the pH between 5 and 10. In particular, a pH between 5.5 to 8.0 and ionic strength of 0.3 or less are preferable to sufficiently ensure an affinity to heparin. Thus, the hepatocyte growth factors adsorbed onto the heparin gel carrier can be recovered by increasing the ionic strength. For example, the recovery can be achieved by using a buffer solution such as sodium phosphate buffer with an addition of inorganic salts such as sodium chloride and ammonium sulfate. The method for the recovery can be a method in which the salt concentration is increased either in a gradient system or a step-wise system. Specifically, the ionic strength to be used is in a range of 0.3 to 3, preferably 0.5 to 2.

The natural-type protein which is a type of hepatocyte growth factor and has a hemopoietic stem cell augmenting activity, which is purified and isolated as described above, can be hydrolyzed using an acid such as hydrochloric acid or proteinases such as pepsin, chymotrypsin and carboxy peptidase; peptide fragments thus obtained are subjected to chromatography such as ion-exchange chromatography to determine their amino acid compositions as well as amino acid sequences.

A process for the analysis of the amino acid composition of the natural-type protein of the present invention, which is a type of hepatocyte growth factor and has the hemopoietic stem cell augmenting activity, will be explained in detail. First, said protein which is a type of hepatocyte growth factor and has the hemopoietic stem cell augmenting activity is hydrolyzed with hydrochloric acid and then the resulting amino acids are converted into corresponding phenylthiocalbamyl derivatives by reaction with phenylisothiocyanate (PITC); the derivatives are then quantitatively analyzed by reversed-phase high performance liquid chromatography (a PITC method).

The amino acid sequence thus obtained can be utilized when the natural-type protein which is a type of hepatocyte growth factor and has the hemopoietic stem cell augmenting activity is produced using genetic recombination techniques.

Next, a process using typical genetic recombination technique for the preparation of the factor, which is a type of hepatocyte growth factor and is used as a hemopoietic stem cell augmenting agent according to the present invention, will be given below.

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Methods for obtaining RNA from human normal fibroblasts include the application of conventional processes such as isolation of polysome, sucrose density gradient centrifugation and electrophoresis. An appropriate method for extracting RNA from the above-mentioned human normal fibroblasts can be selected from various methods including the guanidine thiocyanate-cesium chloride method in which CsCl density gradient centrifugation is carried out after the treatment with guanidine thiocyanate (Chirgwin et al., Biochemistry, 18, 5294, 1979), a method in which a vanadium complex is treated with a surface action agent in the presence of a ribonuclease inhibitor, and then further treated with phenol (Berger et al., Biochemistry, 18, 5143, 1979), a guanidine thiocyanate-hot phenol method, a guanidine thiocyanate-guanidine hydrochloride method, a guanidine thiocyanate-phenol chloroform method and a method in which after the treatment with guanidine thiocyanate, RNA is precipitated by treatment with lithium chloride.

mRNA is isolated from human normal fibroblasts by a conventional method such as the lithium chloride/urea method, guanidine isothiocyanate method and oligo-dT cellulose column method. cDNA is synthesized from mRNA thus obtained using a conventional method such as a method of Gubler et al. (Geen, 25, 236-269, 1983) and a method of H. Okayama et al. (Mol. Cell. Biol., 2, 161, 1982; ibid., 3, 280, 1983). In order to synthesize cDNA from the obtained mRNA, a reverse transcriptase, for example, of avian myeloblastosis virus is principally used, but a method using DNA polymerase can also be used in combination with the partial use of a primer; however, it is convenient to use a commercially available kit for the synthesis or cloning.

The cDNA thus prepared is incorporated in an expression vector, plasmid or baccteriophage and using this recombinant DNA, a cDNA library is constructed with the use of Escherichia coli, according to conventional methods such as the method of Seed (Nature, 329, 840-842, 1987).

To incorporate the above-mentioned cDNA into a vector, a conventional method is applied, using the cohesive ends created using the same restriction enzymes, or synthesized linker site or adapter site are added if necessary, or a homo-polymer is added.

These procedures can be carried out, according to conventional methods (Molecular Cloning, Cold Spring Harbor Laboratory, New York, 1982). The vectors and plasmids into which the cDNA is incorporated are, for example, CDM8, pcDL-SRalpha296, pBR322, pUC18, pUC19 and pUB110 but are not limited to these examples; any vectors and plasmids which are conventionally used for the incorporation of cDNA may be used. Vectors and plasmids which are used to make a cDNA library using E. coli are preferable. In the case where a vector for the incorporation of cDNA is a phage, for example, lambda-gt10 and lambda-gt11 are used but are not limited to these examples; any phages which are conventionally used for the incorporation of cDNA may be used.

In order to introduce a recombinant vector thus obtained into a host, various kinds of conventional methods may be used.

In the case where the vector is a plasmid, an example of a method includes the one in which the vector is introduced into competent cells prepared according to a method of Hanahan et al. (J. Mol. Biol., 166, 557, 1983) in the presence of CaCl<sub>2</sub> or RbCl. In the case where the vector is a phage, an example of a method is the one in which recombinant phage vector is infected into host cells in an appropriate growth phase using an in vitro packaging method or the like.

Examples of the host cells which maintain the cDNA library thus obtained include specifically <u>E. coli</u> MC1061/P3, NM514, NM522, JM101 and C600; however, they are not limited to these examples and any conventional host cells which maintain the cDNA library may be used.

Next, based on the nucleotide sequence of the N-terminal and C terminal of the human hepatocyte growth factor, which is a type of hepatocyte growth factor (Nakamura et al., Nature, 342, 440-443, 1989), an oligonucleotide for use as a probe was synthesized; using this probe in which the oligonucleotide is labeled with <sup>32</sup>P, the targeted cDNA can be obtained by a method such as a colony hybridization method, plaque hybridization method, hybridization-translation assay method and plus-minus method.

More specifically, the DNA in a plaque of the recombinants is immobilized on a filter such as a nylon membrane and then reacted with the labeled probe to select the recombinants having the DNA sequence which binds selectively with this probe.

The above-mentioned probe to be used herein refers to the nucleotide sequence which is complementary to the targeted DNA sequence; the probe can be either DNA or RNA, either chemically synthesized or natural, or else those obtained by the recombinant DNA method; the use of a DNA sequence which is chemically synthesized by means of a known method is general and preferable.

Examples of methods of synthesizing oligonucleotides used herein include conventional methods used for chemical synthesis of nucleic acids, such as the phosphotriester method (Tetrahedron, 34, 3143, 1978; Adv. Carbohydr. Chem. Biochem., 36, 135, 1979; Nucleic Acids Res., 10, 2597, 6553, 1982) and phosphoamidite method (Nature, 310, 105, 1984) and combinations of these methods.

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Next, another method can be used, in which based on the one N-terminal and C-terminal sequence of the above-mentioned human hepatocyte growth factor, which is a type of hepatocyte growth factor (Nakamura et al., Nature, 342, 440-443, 1989), two kinds of primers are synthesized by a DNA synthesizer; in the presence of a DNA polymerase such as Taq DNA polymerase together with each of the primers and cDNA, denaturation of the DNA and then annealing of the primers are allowed to proceed and the elongation reaction of the primers is carried out, for example, using a DNA thermal cycler (a product of Perkin-Elmer Cetus). The resultant product is subjected to electrophoresis and then the targeted cDNA of human hepatocyte growth factor, which is a type of hepatocyte growth factor (HGF) can be prepared according to the conventional method (Molecular Cloning, Cold Spring Harbor Laboratory, New York, 1982).

Furthermore, another method can also be used, in which primers are synthesized using a DNA synthesizer based on the 16 amino acid sequence in one N-terminal nucleotide sequence of the human hepatocyte growth factor, which is a type of hepatocyte growth factor and is obtained by culturing cells such as human fibroblasts, in the presence of a DNA polymerase such as Taq DNA polymerase together with each of the primers and then the denaturation of DNA followed by the annealing of the primers are allowed to proceed and the elongation reaction of the primers is carried out, for example, using a DNA thermal cycler (a product of Perkin-Elmer Cetus). The resultant product is subjected to electrophoresis and then the targeted cDNA of human hepatocyte growth factor, which is a type of hepatocyte growth factor (HGF) can be prepared according to the conventional method. Further, it is observed that when the genes, primers as well as probes of different sources are used, various cDNAs for the human hepatocyte growth factor, which is a type of hepatocyte growth factor, having different nucleotide sequences are accordingly obtained

The recombinant DNA thus obtained is treated with restriction enzymes or the like and the nucleotide sequence of the cDNA is determined according to the conventional method. The nucleotide sequence of the cDNA is obtained by a method such as the Maxam Gilbert method, the dideoxy method of Sanger such as

the dideoxy nucleotide chain termination method (Sanger, Science, 214, 1205, 1981; Methods in Enzymology, 65, 560-580, 1980; Messing, J. et al., Nucleic Acids Res., 9, 309, 1981). Furthermore, the cDNA is synthesized by a primer extension method using the above-mentioned isolated mRNA and a part of the sequenced cDNA and then the recombinant DNA can be obtained as described above.

These methods can be used in appropriate combinations.

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Since cleavage, deletion, addition and binding of DNA chains and furthermore substitutions of nucleotides in the DNA chains according to genetic recombination techniques, can be carried out by conventional procedures, the DNA of the present invention relates not only to the DNA having the nucleotide sequence of SEQ ID NO: 2 in Sequence Listing but also to the above-mentioned altered or modified forms to the extent within the scope of the present invention.

Typical methods for the above-mentioned alteration include a method known as oligonucleotide directed mutagenesis, a method of M. Smith and S. Gillam (Genetic Engineering (J. K. Setlow and A. Hollaender eds, Vol. 3, p.1, 1981) and the methods described in Methods in Enzymology, Vol. 153-155, 1987, Academic Press, CA) and the methods in the references cited in the same.

A particularly preferred method for the alteration and modification of the gene having the nucleotide sequence shown in SEQ ID NO: 2 in Sequence Listing is a method to increase the stability of the targeted protein and to increase biological activity.

Furthermore, there is an altered and modified method in which among the activities of the peptide having the amino acid sequence shown in SEQ ID NO: 2 in Sequence Listing, the activity of proliferation of undifferentiated pluripotent hemopoietic stem cell can be increased.

The cloned cDNA which contains the nucleotide sequence coding for all or part of the amino acid sequence of the protein which is a type of hepatocyte growth factor and has the hemopoietic stem cell augmenting activity, or a substance equivalent in terms of effectiveness is deemed to be a recombinant expression vector in which said coding nucleotide sequence is so incorporated as to be expressed.

In particular, the cDNA which contains the nucleotide sequence coding for all or part of the sequence of the protein or a substance equivalent in terms of effectiveness having the amino acid sequence of SEQ ID NO: 2 in Sequence Listing can be recombined in a suitable expression vector and then the recombinant vector is introduced into an appropriate host cell to form transformants; the transformants thus prepared are cultured and the protein having the targeted hemopoietic stem cell augmenting activity can be obtained by appropriately inducing the expression; the cDNA being thus useful.

In the production of the targeted protein in the host cells using the cDNA which contains the nucleotide sequence coding for all or part of the amino acid sequence of the protein of the present invention, which is a type of hepatocyte growth factor and has the hemopoietic stem cell augmenting activity, or a substance equivalent in terms of effectiveness, the protein can be produced as a mature protein, namely in a form from which signal peptides are removed, or with the above-mentioned signal peptides as they are, or else signal peptides compatible in appropriate host cells or the like are added to and the protein can be produced by secretion from the host cells or the like.

Furthermore, the protein of the present invention, which is a type of hepatocyte growth factor and has the hemopoietic stem cell augmenting activity, or a substance equivalent in terms of effectiveness can be produced as a fused protein or peptide with other recombinant protein or peptide and then the targeted protein is isolated or obtained without isolation by digestion or treatment with enzymes or chemicals from the fused proteins.

In order to effectively express the protein of the present invention, which is a type of hepatocyte growth factor and has the hemopoietic stem cell augmenting activity, or a substance equivalent in terms of effectiveness, the cDNA containing its coding nucleotide sequence can be located at the down stream region which is under the control of a promoter, a ribosome binding site (for example, the SD sequence), a translation initiation site or codon, and then a termination site or codon and a terminator can be located. Namely, in this case, a initiation codon and a stop codon are necessary in the DNA sequence of the gene for its use and are added if necessary using known methods.

As the expression vector to be used herein is not specifically restricted and any vectors which can be autonomously replicable are usable; vectors which contain a replication origin, a selection marker, a promoter, an RNA splicing site and a polyadenylation signal or the like are preferably used.

Examples of the selective marker of the recombinant expression vector include various antibiotic resistant genes, such as ampicilin resistant gene, tetracycline resistant gene and neomycin resistant gene.

Furthermore, examples of the vectors include those derived from various bacteria, those derived from bacteriophages, those derived from animal viruses including insect viruses and mammalian cell viruses, namely, various virus vectors, various plasmid vectors, cosmid vectors, shuttle vectors or the like.

Furthermore, examples of these vectors include <u>E. coli</u>, particularly EK-type plasmid vector, lambda-gt-type phage vectors, vectors derived from <u>Pseudomonas aeruginosa</u>, vectors derived from <u>Bacillus subtilis</u>, vectors derived from yeast, vectors derived from <u>SV40</u>, vectors derived from <u>BPV</u> and vectors derived from retrovirus. Specifically, the examples include pBR322, pUC18, pUB110, pRB15, lambda-gt10, lambda-gt11, SV40 and BPV.

The promoters to be used in the above-mentioned vectors have no particular restriction as long as they can promote expression in the host. Examples of promoters for expression include a tryptophan (trp) promoter, lactose (lac) promoter, a tryptophan/lactose (tac) promoter, a T7 promoter and a lambda-PL promoter which are well known to those skilled in various fields.

Examples of the promoters which are used in vectors for yeast and are typical regulation sequences include promoters relating to the synthesis of enzymes in glycolysis, such as a promoter relating to glycerate-3-phosphokinase, a promoter relating to glyceraldehyde-3-phosphate dehydrogenase, promoters relating to hexokinase, pyruvate decarboxylase, fructose phosphokinase, glucose-6-phosphate isomerase, 3-phosphate isomerase, phosphoglucose isomerase, glucose kinase or the like, or else promoters relating to alcohol dehydrogenase, chytochrome C, acid phosphatase or the like.

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Examples of promoters which are be compatible in mammalian cells include an SV40 early-stage gene or late-stage gene promoter, promoters derived from cytomegalovirus, polyoma virus, adeno virus, bovine papillomavirus or avian sarcoma virus, promoters relating to moloney mouse sarcoma virus LTR, rous sarcoma virus LTR, mouse mammary cancer virus LTR and methallothionein, promoters relating to immunoglobulins, promoters relating to heat shock, promoters relating to dehydrofolic acid, promoters relating to actin and promoters relating to elongation factors.

For insect cells, a promoter relating to polyhydrin derived from nuclear polyhederosis disease virus can be used.

These gene regulating sequences are incorporated into an appropriate vector in combination or in chemically modified form if appropriate and used to construct a vector for the expression of the cDNA which contains the nucleotide sequence coding for all or part of the amino acid sequence of the protein of the present invention, which is a type of hepatocyte growth factor and has the hemopoietic stem cell augmenting activity, or a substance equivalent in terms of effectiveness.

For example, the sequence can contain a translation initiation codon ATG and a termination codon TAA, TGA or TAG as a gene regulating sequence of the cDNA of the present invention; they can be arranged in multiple numbers or in combination with other codons.

In the vectors for the expression of the cDNA of the present invention, a multiple number of the cDNA of the present invention can additionally be incorporated to be expressed.

The vector for the expression of the cDNA which contains the nucleotide sequence coding for all or part of the amino acid sequence of the protein of the present invention, which is a type of hepatocyte growth factor and has the hemopoietic stem cell augmenting activity, or a substance equivalent in terms of effectiveness can be used to obtain a large quantity of cells called transformant which are capable of producing all or part of the amino acid sequence of the protein, which is a type of hepatocyte growth factor and has the hemopoietic stem cell augmenting activity, or a substance equivalent in terms of effectiveness, by a process in which the cDNA is introduced into an appropriate host according to a conventional method which is specifically known for the corresponding host cells and then transform the host cells, and the host cells thus transformed were multiplied by a culture method or the like.

The hosts, particularly cells, herein used can be any cells of <u>E. coli</u> or gram-negative bacteria other than <u>E. coli</u>, such as <u>Pseudomonas</u>, gram-positive bacteria such <u>as Bacillus subtilis</u> and actinomyces or eukaryotic cells such as yeast, mammalian cells, insect cells, plant cells; however, <u>E. coli</u> and mammalian cells such as COS cells and CHO cells are preferably used.

Any methods which are conventionally used in the field of genetic recombination technology can be used as a process for the induction of the gene expression vectors of the present invention into the above-mentioned host cells; examples of methods include a method in which competent cells and the above-mentioned vectors are mixed, a method in which host cells are converted to protoplasts after which the above-mentioned vectors bound to a carrier are introduced into the protoplasts, or a calcium phosphate co-precipitation method, a DEAE dextran method, an electroporation method, an in-vitro packaging method, a virus vector method and a microinjection method.

The transformants thus obtained can be grown under the conditions in which expression of foreign genes therein is suppressed and then the expression of the targeted gene is induced.

The growth or the cultivation of the transformants can be carried out using various kinds of conventional media for cell culture in the same manner as in the case of the above-mentioned human normal fibroblasts. Other growth conditions, culture systems as well as processes for the isolation and purification of the

protein are the same as in the case of human normal fibroblasts. The hemopoietic stem cell augmenting agent of the present invention has an activity to augment undifferentiated pluripotent hemopoietic stem cells and is useful as an active hemopoietic stem cell augmenting agents for the treatment of bone marrow suppression (for example after the use of anti-cancer agents or bone marrow transplantation), as an active hemopoietic stem cell augmenting agent for the treatment of bone marrow malfunctions (such as hypoplastic anemia and osteomyelodysplasia syndrome) or as an active hemopoietic stem cell augmenting agent for the in vitro proliferation of peripheral stem cells and bone marrow stem cells.

Furthermore, since the hemopoietic stem cell augmenting agent eventually promote the growth of not only a variety of blood cells but also progeny of hemopoietic stem cells, osteoclasts, it is applicable as a therapeutic agent for osteoporosis or the like.

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In the present invention, particularly, the recombinant human normal fibroblast protein which is a type of hepatocyte growth factor and has the amino acid sequence of SEQ ID NO: 2 in Sequence Listing, or a substance equivalent in terms of effectiveness, is preferably used as a hemopoietic stem cell augmenting agent, exhibits an augmenting activity on pluripotent hemopoietic stem cells, is an active hemopoietic stem cell augmenting agent to restore bone marrow growth (for example after the use of anti-cancer agents or bone marrow transplantation), is an active hemopoietic stem cell augmenting agent for the treatment of bone marrow malfunctions (such as hypoplastic anemia and osteomyelodysplasia syndrome) or is an active hemopoietic stem cell augmenting agent for the in vitro proliferation of peripheral blood stem cells and bone marrow stem cells. The protein is derived from human normal fibroblasts and accordingly is a relatively normal-type, which is considered to be preferable.

Furthermore, a corresponding natural-type protein obtained from human normal fibroblasts is considered to be preferable as a hemopoietic stem cell augmenting agent and is useful in the above-mentioned utilization.

In the cases where the hemopoietic stem cell augmenting agent of the present invention is used for the above-mentioned uses according to the present invention, it can be administered orally or parenterally as it is or as a pharmaceutical composition mixed with a pharmaceutically acceptable vehicle, an excipient or the like, which is known per se.

In order to maintain an active component in the hemopoietic stem cell augmenting agent of the present invention, amino acids such as arginine, lysine, glycine, leucine, phenylalanine and aspartic acid, saccharides such as glucose, sucrose, mannitol and mannit, polysaccharides or protein hydrolysates such as sugar alcohol, gelatin, collagen, dextran, pullulan, heparin, chondroitin sulfate and cellulose, inorganic acids such as hydrochloric acid or organic acids such as acetic acid, inorganic bases such as sodium hydroxide or inorganic bases such as amines or the like can be added when appropriate.

Dosage forms of the preparations for oral administration are specifically tablets, pills, capsules, syrups, emulsions, suspensions or the like. The preparations in such forms can be manufactured by the methods known per se and contain vehicles or excipients which are conventionally in the pharmaceutical field. Examples of vehicles and excipients for tablets include lactose, starch, sucrose and magnesium stearate.

Dosage forms of preparations for parenteral administration are, for example, ointment, injectable preparations, wet compressing agents, endermic liniments, agents, inhalants, suppositories, percutaneous vapors or the like. The injectable preparations can be prepared according by means known per se such that the hepatocyte growth factor of the present invention is dissolved, suspended or emulsified in a sterilized aqueous or oily solution which is conventionally used for injectable solutions. Examples of the aqueous solutions for injectable solutions include physiological saline and glucose solutions and examples of the oily solutions include sesame oil and soybean oil; in each case solubilizers may be used in combination. Suppositories used for rectal administration are prepared by means known per se such that the hemopoietic stem cell augmenting agent of the present invention is mixed with a conventional base for suppositories and the mixture is formulated.

The effective quantity and number of doses of the hemopoietic stem cell augmenting agent of the present invention depends on such factors as mode of administration, age and body weight of patient and property and severity of the particular disease to which the therapy is applied, however, in general, the daily dose for an adult is between 0.01 and 100 mg, preferably between 0.1 and 10 mg in one dose or in divided doses.

The novel protein of the present invention, which is a type of hepatocyte growth factor and has a hemopoietic stem cell augmenting activity, or a substance equivalen in terms of effectiveness can produce antibodies in such a way that the protein or its equivalent alone or in a form conjugated with a vehicle such as bovine serum albumin, egg serum albumin, thioglobulin or hemocyanin (KLH) by means of, for example, carbodiimide, glutaraldehyde, mixed acid anhydrates, homo-difunctional or hetero-difunctional reagents (e.g. maleidobenzyl-N-hydroxysuccinimide ester (MBS) or the like is injected to an animal for immunization.

Furthermore, spleen cells obtained from the animal such as a mouse thus immunized and myeloma cells are fused according to the conventional method and thus hybridoma cells to produce monoclonal antibodies can be obtained.

The antibodies thus obtained can be made into a reagent for measurement by binding onto a solid phase, or by binding with, for example, enzymes, co-enzymes, fluorescent substances, coloring agents such as dye, radio-active labels or paramagnetic metals. Examples of standard methods for immunoassay are an enzyme-linked immunoassay (ELISA), a radio immunoassay (RIA) and a sandwich immunoassay.

These assay are carried out according to D. Catty, "Antibodies - Vol. I & Vol. II, A practical approach", IRL Press.

10 EXAMPLES

The invention is further illustrated by the following examples but the invention is not limited to these examples.

Example 1

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Preparation of human hepatocyte growth factor derived from normal fibroblasts:

Using a 30-I culture bath, cells of human normal fibroblast DIP2 (Kobayashi, S. et al., in The clinical potential of interferons, ed. Kono, R. and Vilcek, J., University of Tokyo Press, Tokyo, 1982) were cultured on 10% FBS-MEM medium-0.3% beads (Cytodex-1, Pharmacia) at 37°C for 5 days with stirring. When growth reached the confluent state, the medium was changed to a fresh MEM medium and protein production was induced by the addition of poly I/C (10 micrograms/ml). The incubation was continued at 37°C for another 4 days and then 20 liters of culture supernatant were recovered.

The culture supernatant was adsorbed on a "Blue Sepharose Column" (1 liter gel carrier; Pharmacia) equilibrated with a 20 mM tris-HCl buffer solution (pH 8.0) and then the column was washed with the same buffer used for the equilibration. Subsequently, proteins were eluted using a 0-3 M linear NaCl concentration gradient.

One liter of an active fraction thus obtained was desalted by gel filtration and then the desalted fraction was adsorbed on a heparin-Sepharose column (gel carrier 100 ml; Pharmacia) equilibrated with a 20 mM tris-HCl buffer solution (pH 8.0) and the column was washed with the same buffer used for the equlibration. Subsequently, proteins were eluted using a 0-3 M NaCl linear concentration gradient.

50 ml of an active fraction thus obtained was desalted by gel filtration and then purified by reversed-phase high performance liquid chromatography. Using a "Vydac 218TP510 Column" (1.0 x 25 cm; Separation Group), elution was carried out using a 0-50% acetonitrile linear concentration gradient in a 0.1% aqueous trifluoroacetic acid solution. Results are shown in Figure 9.

A fraction of the active peak was subjected to SDS polyacrylamide electrophoresis under reducing conditions (Laemmli U. K., Nature, 227, 680-685, 1970); a single band at a molecular weight of about 60K was obtained.

Example 2

N-terminal amino acid sequence and amino acid composition:

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The purified protein obtained in Example 1, which is a natural-type human hepatocyte growth factor derived from fibroblasts, (referred to as natural-type human HGF hereinafter) was placed in an amino acid sequencer (Applied Biosystems 477A Protein Sequenser); as a result, the sequence of 16 amino acids at the N-terminal was determined as shown in SEQ ID NO: 1 in Sequence Listing. Judging from the homology in sequence of this N-terminal sequence with that disclosed by Nakamura et al. (Nature, 342, 440-443, 1989), this sequence was revealed to be the N-terminal of the beta chain.

25 microliters of conc. hydrochloric acid containing 0.4% thioglycolic acid was added at a concentration of 4 micrograms/25 microliters; after hydrolysis at 110°C for 22 hours in a tube sealed in vacuum, the hydrochloric acid was dried under reduced pressure. The resultant product was dissolved in distilled water and subjected to amino acid analysis using an amino acid analyzer (Hitachi 835-type amino acid analyzer 9). Results are shown in Table 1.

Table 1

Amino acid	Mol %	Amino acid	Mol %
Asp + Asn	12.85	Met	2.00
Thr	5.76	lle	5.27
Ser	6.31	Leu	5.87
Glu + Gln	9.67	Tyr	4.73
Pro	5.95	Phe	2.72
Gly	9.91	Lys	6.45
Ala	3.86	His	3.45
1/2 Cys	3.31	Trp	1.21
Val	4.83	Arg	5.86

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Example 3

Cloning of human hepatocyte growth factor cDNA derived from normal fibroblasts:

Since cDNA of human hepatocyte growth factor had been cloned and sequenced by Nakamura et al. (Nature, 342, 440-443, 1989), a primer was synthesized based on the sequence; after replicating cDNA by the polymerase chain reaction (PCR) method, the cDNA can be cloned in, for example, an expression vector.

(1) Isolation of human normal fibroblasts mRNA:

RNA was prepared from human normal fibroblasts MRC5 (RCB211, available at Riken Cell Bank, Japan), which was cultured in the same manner as in Example 1, according to a lithium chloride/urea method (Auffray et al, Eur. J. Biochem., 107, 303-314, 1980). RNA thus obtained was dissolved in a 10 mM tris-HCl buffer solution (pH 7.5) containing 1 mM EDTA (referred to as TE hereinafter) and after heating the solution at 70°C for 5 minutes, an equal amount of TE containing 1 M LiCl was added to the solution. The resultant RNA solution was applied on an oligo-dT cellulose column which had been equilibrated with TE containing 0.5 M LiCl and the column was washed with the same buffer solution. The column was further washed with TE containing 0.3 M LiCl and then adsorbed poly (A) RNA was eluted with 2 mM EDTA (pH 7.0) containing 0.01% SDS.

(2) Construction of cDNA library derived from human normal fibroblasts:

cDNA was prepared using 4 micrograms of poly (A) RNA obtained as described in (1) above, according to the method of Gubler et al. (Geen, 25, 236-269, 1983).

This cDNA was inserted in an expression vector CDM8 using T4 DNA ligase according to the method of Seed (Nature, 329, 840-842, 1987). Using this recombinant DNA, E. coli MC1061/P3 was transformed to obtain cDNA libraries. The titration revealed that this cDNA library consisted of 200 thousand transformants. Plasmid DNAs were isolated from these transformants according to the conventional method (Molecular Cloning, Cold Spring Harbor Laboratory, New York, 1982).

(3) Isolation of human hepatocyte growth factor cDNA derived from normal fibroblasts:

Appropriate sequences are selected from the amino acid sequence of the human hepatocyte growth factor: for example, a nucleotide sequence of the N-terminal or C-terminal is selected and two kinds of primers are synthesized base on the selected sequence by a DNA synthesizer; after amplifying cDNA by the polymerase chain reaction (PCR) method, the products are cloned in an expression vector or the like.

Two kinds of primers having the following sequences were synthesized by a DNA synthesizer based on the human hepatocyte growth factor sequences of the N-terminal and C-terminal of a human hepatocyte growth factor derived from the liver, known as a human hepatocyte growth factor (Nakamura et al., Nature, 342, 440-443, 1989).

5' ATGTGGGTGACCAAAC 3'

#### 5' CTATGACTGTGGTACC 3'

20 pmol of each primer and 1 microgram of the plasmid DNA obtained in the above-mentioned (2) were placed in a micro-centrifuge tube and then various reagents were added to the tube to make a mixture of a total volume of 100 microliters containing 20 mM tris-HCl buffer (pH 8.3), 1.5 mM MgCl<sub>2</sub>, 25 mM KCl, 100 micrograms/ml gelatin, and 50 microM each of dNTP, and 4 unit Taq DNA polymerase. The resulting mixture was subjected to reactions of 40 cycles using a DNA thermal cycler (Perkin-Elmer Cetus) in which the conditions for DNA denaturation were at 94°C for 1 minute, the conditions for primer annealing were at 50°C for 3 minutes and the conditions for primer extension were at 72°C for 3 minutes. The reaction product was subjected to a 1% agarose-gel electrophoresis and a human hepatocyte growth factor cDNA derived from normal fibroblasts, having a size of about 2.2 kb, was prepared according to the conventional method (Molecular Cloning, Cold Spring Harbor Laboratory, New York, 1982).

Alternatively, the human hepatocyte growth factor derived from normal fibroblasts can be obtained by a colony hybridization method in which an oligomer is synthesized by a DNA synthesizer based on the amino acid sequence (having 16 amino acids) of the N-terminal of the human hepatocyte growth factor derived from human normal fibroblasts prepared as described in Example 1.

#### (4) Preparation of expression vector:

An expression vector CDM8 (Seed, Nature, 329, 840-842, 1987) was digested with restriction enzyme HindIII, and blunt ends were made with T4 DNA polymerase and an EcoRI linker was herein ligated using T4 DNA ligase. Then, the resultant fragment was digested with restriction enzyme PstI and blunt ends were made using T4 DNA polymerase. A KpnI linker was ligated using T4 DNA ligase and then the resultant fragment was digested with restriction enzymes EcoRI and KpnI. The resultant product was subjected to 1% agarose-gel electrophoresis and a DNA fragment of about 0.36 kb was prepared according to the conventional method. On the other hand, pcDL-SRalpha296 (Takebe et al., Mol. Cell. Biol., 8, 446-472, 1988) was digested with restriction enzymes EcoRI and KpnI and a DNA fragment of about 3.4 kb was subsequently purified by agarose-gel electrophoresis; the vector thus obtained was ligated with the DNA fragment of about 0.36 kb obtained as described above, using T4 DNA ligase. Using this product, E. coli was transformed by the conventional method and a plasmid DNA was obtained from the resultant transformant according to the conventional method (Molecular Cloning, Cold Spring Harbor Laboratory, New York, 1982) and then a targeted expression vector, pSRalphaBX was obtained.

Figure 1 illustrates the construction of expression vector pSRalphaBX.

This plasmid DNA was digested with restriction enzyme BstXI and the reaction solution was subjected to 1% agarose-gel electrophoresis, thereby a DNA fragment of 3.4 kb both ends having a restriction enzyme BstXI cleavage site was isolated and purified.

(5) Cloning of human hepatocyte growth factor cDNA derived from normal fibroblasts into expression vector and determination of nucleotide sequece:

The 2.2 kb cDNA fragment of the human hepatocyte growth factor derived from normal fibroblasts which was obtained as described above in (3) was phosphorylated using T4 DNA kinase according to the conventional method (Molecular Cloning, Cold Spring Harbor Laboratory, New York, 1982) and then ligated with a BstXI linker (N408-18; Invitrogen) using T4 ligase. Subsequently, this reaction mixture was subjected to 1% agarose-gel electrophoresis, whereby a DNA fragment of 2.2 kb ligated with BstXI linker was isolated and purified. This DNA fragment was ligated using T4 ligase with the DNA fragment of 3.4 kb obtained in the above-mentioned (4), in which both ends were digested with restriction enzyme BstXI. Using this product E. coli was transformed and a plasmid DNA was prepared according to the conventional method from the transformant thus obtained. Subsequently, the plasmid carrying the cDNA fragment of human hepatocyte growth factor derived from normal fibroblasts was confirmed by digesting this plasmid DNA with restriction enzyme BamHI (said plasmid is referred to as pSRαFDF-1 or pSRalphaFDF-1 hereinafter) and then the nucleotide sequence of the cDNA of hepatocyte growth factor derived from normal fibroblasts was determined by the dideoxy method (Prober et al., Science, 238, 336-341, 1987) using the Genesis 2000 DNA analysis system (DuPont) (Figure 2).

Figure 3 illustrates the construction of human HGF expression vector pSR $\alpha$ FDF-1 for expression in mammalian cells.

(6) Expression of human hepatocyte growth factor gene isolated from normal fibroblasts in monkey COS cells:

10 micrograms of pSRalphaFDF-1 obtained as described in (5) above was added to 4 ml of an RPMI164 medium containing 40 mM tris-HCl buffer (pH 7.5), 400 microgram/ml DEAE dextran (Pharmacia) and 100 microM chloroquine(Sigma). On the other hand, COS-1 cells (ATCC CRL-1650), which had been grown to confluent in an RPMI1640 medium (Gibco) supplemented with 10% fetal calf serum (Gibco), was washed once with PBS and then 4 ml of the above-mentioned DNA mixture was added to the cells, after which the incubation was carried out at 37°C in an atmosphere of 5% CO<sub>2</sub>. After 4 hours, cells were washed with PBS and then cultured in 20 ml of RPMI1640 medium at 37°C under 5% CO<sub>2</sub> for 4 days. The human hepatocyte growth factor activity in the culture supernatant was 340 units/ml when measured using the growth of NFS60 cells as the index. On the other hand, in the culture supernatant which was obtained in the same manner except that a vector in which the cDNA of said human hepatocyte growth factor activity was observed.

(7) Expression of human hepatocyte growth factor gene derived from normal fibroblasts in Chinese hamster CHO cells:

The CHO clone DUKXB11 cells (provided by Dr. Chasin of Columbia University), a dihydrofolic acid reductase (DHFR) defective strain of Chinese hamster CHO cells, were cultured overnight in an alpha-MEM (Gibco) medium supplemented with 10% fetal calf serum and nucleic acids at a concentration of 1 x 10<sup>5</sup> per well of a 12-well plate.

1 microgram of pSRalphaFDF-1 and 0.1 microgram of pAdD26SV(A)-3 (Scahill, Proc. Natl. Acad. Sci. USA, 80, 4654-4658;, 1983) were mixed and introduced into the above-mentioned CHO cells using a transfection kit by Pharmacia and the cells were cultured for 18 hours. Then, the cell culture was diluted 20 times and incubated for 10 days in an alpha-MEM (Gibco) without nucleic acid and supplemented with 10% fetal calf serum and transformants were obtained.

From the cell strains thus obtained, strains which exhibited high hepatocyte growth factor activity in the culture supernatant were selected and the cells of the selected strains were cultured in an alpha-MEM (Gibco) without nucleic acid and supplemented with 50 nM methotrexate and 10% fetal calf serum; and thus a clone having a high hepatocyte growth factor productivity was obtained and named CHO-6-23-2. The production of human hepatocyte growth factor derived from normal fibroblasts by these cloned cells was 3500 units/ml/2days when measured using NFS60 cells proliferation assay as the index.

#### Example 4

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A recombinant human hepatocyte growth factor derived from normal fibroblasts was purified from the culture supernatant of the human hepatocyte growth factor-producing monkey COS-1 cells obtained in Example 3 (6).

(1) Salting-out with ammonium sulfate:

5265 g of ammonium sulfate were gradually added to 13.5 liters of the COS-1 cell culture supernatant and dissolved and then the solution was allowed to stand at 4°Covernight. The precipitate was collected by centrifugation at 6500 rpm for 20 minutes and was dissolved in a 20 mM tris-HCl buffer solution (pH 8.0). The solution was thoroughly dialyzed against the same buffer solution to prepare a concentrated solution by ammonium sulfate.

(2) Anion exchange chromatography:

The concentrated, ammonium sulfate treated solution obtained in the above-mentioned (1) was added to 10 ml of DEAE Sephacel (Pharmacia) which had been equilibrated with a 20 mM tris-HCl buffer solution (pH 8.0). Unadsorbed substances were washed with a 20 mM Tris-HCl buffer solution (pH 8.0) and then adsorbed substances were eluted by adding in order 100 ml portions of 20 mM tris-HCl buffer solution (pH 8.9) containing NaCl in concentrations of 0.05 M, 0.3 M or 0.5 M. The chromatogram pattern is illustrated in Figure 4. Fractions showing the NFS60 cell proliferating activity were collected to make a DEAE Sephacel eluate.

#### (3) Heparin-Sepharose CL-6B chromatography:

The DEAE Sephacel eluate was added to 2ml of heparin-Sepharose CL-6B (Pharmacia) which had been equilibrated with a 20 mM tris-HCl buffer solution containing 0.3 M NaCl (pH 8.0). After thoroughly washing with a 20 mM tris-HCl buffer solution containing 0.3 M and then with the same buffer containing 0.5 M NaCl, elution was carried out with a 20 mM tris-HCl buffer solution containing 1 M NaCl (pH 8.0). The chromatogram pattern is illustrated in Figure 5. Fractions having the NFA60 cell proliferating activity were collected to make a heparin eluate.

#### (4) Zinc chelate affinity chromatography:

0.3 ml of chelating Sepharose 6B (Pharmacia) was packed in a column, a 0.5% aqueous zinc chloride solution was added to the column and the column was washed with a 20 mM tris-HCl solution (pH 8.0) to prepare a column for zinc chelate affinity chromatography. 12 ml of a heparin eluate was added to the column and the column was washed with a 20 mM tris-HCl buffer solution containing 1 M NaCl (pH 8.0). Furthermore, the column was washed with a 20 mM tris-HCl buffer solution containing 50 mM NH<sub>4</sub>Cl (pH 8.0) and then elution was carried out with a 20 mM tris-HCl buffer solution containing 50 mM imidazole and 0.5 M NaCl. The elution pattern is illustrated in Figure 6. Fractions having the NFS60 cell proliferating activity were collected to make a zinc eluate. The yield of the purified recombinant-type hepatocyte growth factor was about 750 micrograms and recovery of the activity from the ammonium sulfate treated concentrate was about 44 %.

#### (5) SDS polyacrylamide gel electrophoresis:

The recombinant-type hepatocyte growth factor prepared by the process described above was applied to SDS-polyacrylamide electrophoresis (4-20% gel) under non-reducing conditions. The recombinant-type hepatocyte growth factor exhibited a single band at a molecular weight between 66,000 and 85,000 under non-reducing conditions. Results are shown in Figure 7.

#### 80 Example 5

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A recombinant-type human hepatocyte growth factor derived from normal fibroblasts was purified from a culture supernatant of human hepatocyte growth factor-producing chinese hamster CHO recombinant cell strain CHO-6-23-2 obtained in Example 3 (7).

#### (1) Cation exchange chromatography:

25 ml of a CHO-6-23-2 cell culture fluid was thoroughly dialyzed with a 20 mM tris-HCl buffer solution (pH 6.8) and the dialyzate was added to 0.5 ml of CM-Sephadex (Pharmacia) which had been equilibrated with a 20 mM tris-HCl buffer solution (pH 6.8). After washing with a 20 mM Tris-HCl buffer solution (pH 6.8), the elution was carried out with a 20 mM tris-HCl buffer solution (pH 6.8) containing 0.5 M NaCl. Fractions showing the NFS60 cell proliferating activity were collected to make a CM-Sephadex eluate.

#### (2) Heparin-Sepharose CL-6B chromatography:

The CM-Sephadex eluate was added to 0.1 ml of heparin-Sepharose CL-6B which had been equilibrated with a 20 mM tris-HCl buffer solution containing 0.5 M NaCl (pH 8.0). After washing with a 20 mM tris-HCl buffer solution containing 0.5 M Nacl, elution was carried out with a 20 mM tris-HCl buffer solution containing 1 M NaCl (pH 8.0). Fractions having the NFA60 cell proliferating activity were collected to make a heparin eluate fraction.

#### (3) Zinc chelate affinity chromatography:

0.1 ml of chelating Sepharose 6B (Pharmacia) was packed into a column, a 0.5% aqueous zinc chloride solution was added to the column and the column was washed with a 20 mM tris-HCl solution (pH 8.0) to prepare a column for zinc chelate affinity choromatography. The heparin eluate was added to the column and the column was washed with a 20 mM tris-HCl buffer solution containing 1 M NaCl (pH 8.0). Furthermore, the column was washed with a 20 mm tris-HCl buffer solution containing 50 mM NH<sub>4</sub>Cl (pH

8.0), and then elution was carried out with a 20 mM tris-HCl buffer solution containing 50 mM imidazole and 0.5 M NaCl. Fractions having the NFS60 cell proliferating activity were collected to make a zinc eluate.

(4) SDS polyacrylamide gel electrophoresis:

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The recombinant-type hepatocyte growth factor prepared by the process described above was applied to SDS-polyacrylamide electrophoresis (4-20% gel) under non-reducing conditions; the gel was stained by a silver staining method. The recombinant-type hepatocyte growth factor exhibited a single band at a molecular weight between 66,000 and 85,000 under non-reducing conditions.

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Example 6

Measurement of proliferating activity on murine immature myeloblasts (NFS60):

Cell proliferation was measured as follows according to the MTT Assay method (T. Mosman, J. Immunological Methods, 65, 55-63, 1983):.

A 50 microliter aliquot of a culture medium solution (10% FBS-RPM1640) was placed into each well of a 96-well microplate, a 50 microliter aliquot of a solution containing the natural-type hepatocyte growth factor (natural-type human HGF) derived from human normal fibroblasts purified in Example 1 was added to the medium and then a two-step dilution was carried out. A 50 microliter aliquot of 2 x 10<sup>5</sup> cells/ml suspension of NFS60 cells was placed in each well and then the plate was cultured at 37°C for 2 days in a CO<sub>2</sub> gas incubator.

Subsequently, a 10 microliter aliquot of MTT reagent [prepared by dissolving 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazonium bromide in PBS at 5 mg/ml] was added to each well and the plate was cultured at 37°C for 5 hours in a CO<sub>2</sub> incubator. 150 microliters of 0.04 N hydrochloric acid-added isopropanol was herein added to extract pigment and optical density at 590 nm was determined using an immuno reader.

The natural-type hepatocyte growth factor derived from human normal fibroblasts, purified in Example 1 (a natural-type human HGF), showed activity at a dilution rate where the OD 590 value was 50% of that for the maximum response.

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Example 7

Determination of DNA synthesis stimulating activity on mature rat primary culture hepatocytes:

Mature rat hepatocytes were isolated and purified by the method of Seglen (Seglen, P. O., Methods in Cell Biology, 13, 29-83, 1976).

Freshly prepared hepatocytes (1 x 10<sup>5</sup>) were added to a medium of the following composition and the total volume was made into 1 ml; a medium containing MEM (Gibco), 100 mM insulin (Sigma), 50 micrograms/ml gentamycin (Sigma), 5% calf serum (Gibco) and HGF was dispensed in a collagen-coated 35-mm plastic petri-dish (Falcon). Incubation was carried out in an incubator in an atmosphere of 7% CO<sub>2</sub> and 90% humidity at 37°C for 4 hours. After incubation, the medium was changed with a fresh serum-free MEM medium containing 5 microCi/ml[³H] thymidine. The incubation was continued for another 45 hours under the same conditions as described above. The cultured petri-dish was washed 6 times with 0.9% NaCl. The cells were dissolved in 1.5 ml of 0.33 N NaOH and then the resultant solution was transferred into a test tube in ice water.

0.5 ml of 40% trichloroacetic acid dissolved in 1.2 N hydrochloric acid was added to the solution and the resultant precipitate was isolated by centrifugation at 2,000 rpm for 10 minutes. The precipitate thus obtained was dissolved in 0.5 ml of 0.33 N NaOH and a 0.3 ml portion of the solution was placed into a scintillation vial and then 0.5 ml of Aquasol (New England Nuclear) and 0.4 ml of 40% trichloroacetic acid in a 1.2 N hydrochloric acid solution was added to this vial. Uptake of [3H] thymidine incorporation was measured by a scintillation counter. Hepatocyte growth factor derived human normal fibroblasts purified in Example 1 (natural-type human HGF) exhibited the activity.

#### Example 8

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Measuremet of growth stimulating activity on hemopoietic stem cells using human normal bone marrow cells:

2 to 3 ml of heparin-added normal human bone marrow blood was sampled and allowed to stand in the presence of silica at 37°C for 30 minutes and then non-phagocytic monocytes were isolated by Ficoll-Paque (Pharmacia) gravity centrifugation. After washing, adhesive cells were removed using a 10-cm plastic petridish and the remaining non-phagocytic, non-adhesive monocytes (NPNAMNC) were allowed to suspend in an alpha-medium (Flow Labs).

Culture was carried out by a methyl cellulose method which is a modified method of Iscove et al. (Iscove, N. N. et al., J. Cell Physiol., 83, 309-320, 1974). The above-mentioned NPNAMNC (4 x  $10^4$  cells) were added to a serum free medium having the following composition and the total volume of the medium was made into 1 ml. The medium contained alpha-medium, 0.8% methyl cellulose (Shin-etsu Kagaku), 0.1% crystallized deionized bovine serum albumin (Sigma), 300 micrograms/ml Fe-saturated human transferrin (Sigma), 40 micrograms/ml soy bean lecithin (Sigma), 24 microgram/ml cholesterol (Nakarai Chemical),  $5 \times 10^{-5} \,\mathrm{M}$  2-mercaptoethanol and was dispensed into a 35-mm Lux culture dish (Miles Labs) with samples added to the medium. Incubation was carried out in an incubator in an atmosphere of  $5\% \,\mathrm{CO}_2$  and with 100 % humidity at  $37^{\circ}\mathrm{C}$  for 18 days. Then colonies were observed under an inverted microscope and the number of colonies was counted.

Results are shown in Table 2.

Table 2

25	Hemopoietic st	em cells gr	owth stimulating acti m	vity of natural-type arrow cells	human HGF on hu	man normal bone
	Natural-type HGF added		Number o	of colony formed/4 >	x 10⁴ NPNAMNC	
30		Blast	CFU-GM	CFU-G	Macrophage	Total
	None 1ng/ml	10 6	19 26	5 7	6 1	40 40
35	10ng/ml 10²ng/ml	17 13	38 38	9 9	1 4	65 64
	10 <sup>3</sup> ng/ml	15	30	4	5	54

#### Example 9

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Measurement of growth promoting activity on hemopoietic stem cells using murine bone marrow cells:

150 mg/kg of 5-fluorouracil was injected intravenously to a BDF1 female mouse and after 48 hours bone marrow cells were taken from the thigh bone.

Culture was carried out by a methyl cellulose method which is a modified method of Iscove et al. 5 x 10<sup>4</sup> bone marrow cells were added to a serum-free medium having the following composition and the total volume of the culture was made into 1 ml. The medium contained alpha-medium, 0.9% methyl cellulose, 1% crystallized deionized bovine serum albumin, 300 micrograms/ml Fe-saturated human transferrin, 160 micrograms/ml soy bean lectin (Sigma), 96 micrograms/ml cholesterol (Nakarai Chemical), 10<sup>-4</sup> M 2-mercaptoethanol and samples or various hemopoietic factors added; the medium thus prepared was dispensed into 35-mm Lux culture dishes. Each hemopoietic factors was added at the following concentration: 200 u/ml for rmulL-3 (Cosmo Bio Co., Ltd.) and 20 u/ml for rmulL-7 (Cosmo Bio Co., Ltd.).

Incubation was carried out in an incubator in an atmosphere of 5% CO<sub>2</sub> and 100 % humidity at 37°C. After incubation for 17 days, colonies were observed under an inverted microscope and the number of colonies was counted.

Results for the natural-type hepatocyte growth factor derived from human normal fibroblasts are shown in

Table 3

Natural-typ HGF added		Colonies /5	x 10⁴ 5FU-resistant	bone marrow cells	
	Blast	CFU-GM	CFU-G	Macrophage	Total
None	0	0	0	0	0
10 <sup>2</sup> ng/ml	0	0	0	0	0
+ IL-3	0	1	0	0	1
+ IL-3 + IL-7	8	8	5	2	23
10 <sup>3</sup> ng/ml	12	14	0	6	32

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#### Example 10

Measurement of hepatocyte growth promoting activity of recombinant-type and natural-type HGF derived from human normal fibroblasts:

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Hepatoparenchyma cells were isolated from a 4-week-old Wister rat by a collagenase flux method. The hepatoparenchyma cells thus obtained were suspended in an Williams E medium containing 5% fetal calf serum, 1 x  $10^{-9}$  M insulin and 1 x  $10^{-9}$  dexamethasone at a concentration of  $10^5$  cells/ml. 0.5 ml portions of the cell suspension thus prepared were dispensed into a collagen-coated 23-well multiplate and the plate was incubated at  $37^{\circ}$ C for 20 hours in the presence of 5%  $CO_2$ . Subsequently, the medium was exchanged by a fresh Williams E medium containing 1 x  $10^{-9}$  insulin and 1 x  $10^{-9}$  M dexamethasone and at the same time a specified amount of samples were added. Incubation was continued for another 23 hours and then 0.5 microCi per well of  $^{125}$  I deoxyuridine was added and incubation was continued for another 7 hours. The cells thus obtained were washed twice with PBS and then treated with a cold 10% trichloroacetic acid aqueous solution. The cells were solubilized with 0.5 ml per well of 1N NaOH and the radioactivity of the solution was measured by a gamma counter. A portion of the sample after the radioactivity measurement was taken and measured for the amount of protein by the Lawry method. Radioactivity incorporated in the hepatoparenchyma cells when various samples were added was measured and was converted to per 1 microgram of hepatoparenchyma cell protein to give DNA synthesis activity (cpm/microgram protein).

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Results are shown in Table 4.

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Table 4

Hepatocyte proliferating activity of recombinant-type and

natural-type HGF derived from human normal fibroblasts

10	Cond	centration	DNA synthesis hepatoparenchy (cpm/microgram ce	ma cells
15	Natural-type human HGF	100 ng/ml 10 ng/ml 1 ng/ml 100 pg/ml 10 pg/ml	$   \begin{array}{ccccccccccccccccccccccccccccccccccc$	5 0 0
20 25	Recombinant-type human HGF	100 ng/ml 10 ng/ml 1 ng/ml 100 pg/ml	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	1 6 5
	Epithelial cell growth factor	50 ng/m] 10 ng/m] 2 ng/m]	. 58 <u>±</u> 5.	9
30	Insulin	100 nM 10 nM 1 nM	54 ± 3. 43 ± 1. 32 ± 4.	5
35	Insulin + Epithelial cell growth factor	100 nM 20 ng/ml	100 ± 3.	5
40	Natural-type Human HGF + Insulin	50 ng/ml	137 ± 4.	4
45	Recombinant-type human HGF + Insulin	50 ng/ml 100 nM	118 ± 3.	5
70	Natural-type human HGF + Epithelial cell	50 ng/ml 20 ng/ml	_	9

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Table 4 (continued)

Hepatocyte proliferating activity of recombinant-type

and natural-type HGF derived from human normal fibroblasts

10		Concer	ntration	DNA synthesis in hepatoparenchyma (cpm/microgram cell	
15	Recombinant-type human HGF +	50	ng/ml	110 ± 5.1	
	Epithelial cel growth factor	20	ng/ml		
20	Natural-type human HGF +	50	ng/ml	153 ± 7.5	
	Insulin +	100	nM		
	Epithelial ce growth factor	20	ng/ml		
25					
	Recombinant-type human HGF +	50	ng/ml	138 ±10.7	
	Insulin +	100	nM		
30	Epithelial cell growth factor	20	ng/ml		

Both natural-type HGF which was purified in Example 1 and derived from human normal fibroblasts and recombinant type HGF which was purified in Example 4 and derived from COS cells exhibited DNA synthesis activity at 10 ng/ml and in the presence of insulin and/or epithelial cell growth factor, DNA synthesis activity on hepatoparenchyma cells was enhanced.

#### Example 11

40 Colony formation stimulating activity of recombinant-type and natural-type HGFs derived from human normal fibroblasts using normal murine bone marrow cells:

Bone marrow cells were taken from the thigh bone of a BDF1 mouse and according to the conventional method (Metcalf Clonal Culture of Hemopoietic Cells: Techniques and Applications, Elsevier, Amsterdam), 2 x  $10^4$  of the bone marrow cells were suspended in 1 ml of alpha-MEM medium containing 1 x  $10^{-4}$  M 2-mercaptoethanol, 20% fetal calf serum and HGF sample in various concentrations; incubation was carried out at  $37^{\circ}$ C for 7 days in the presence of 5%  $CO_2$  and then colonies formed were observed under an inverted microscope and counted.

Results are shown in Table 5.

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Table 5
Colony assay using untreated bone marrow cells

10	Natural-type HGF derived from human normal fibroblasts (Unit/ml)	Number	of GM	clusters
15	0 25 100 400 1600	$\begin{array}{ccc} 9.8 & \pm \\ 7.0 & \pm \end{array}$		٠.
20	Recombinant-type HGF derived from human normal fibroblasts (Unit/ml)	Number	of GM	clusters
25 30	0 25 100 400 1600	2.3 ± 9.8 ± 20.0 ± 15.0 ± 15.8 ±	3.3 5.5 5.5 3.6 1.3	

Example 12

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Measurement of NFS60 cell proliferating activity of hepatocyte growth factor derived from human placenta:

NFS60 cell proliferating activity of hepatocyte growth factor derived from human placenta (Becton Dickinson Labware) and natural-type hepatocyte growth factor derived from human normal fibroblasts, which was purified in Example 1, was measured by the method described in Example 6. Results are shown in Table 8.

In the same manner, NFS60 cell proliferating activity of the recombinant-type hepatocyte growth factor derived from human normal hepatocytes (Nature, 342, 440-443, 1989) and a recombinant-type hepatocyte growth factor derived from fibroblast M426 of human fetal lung (Proc. Natl. Acad. Sci. USA, 88, 415-419, 1991) can be measured by the method described in Example 6.

### POSSIBILITY OF INDUSTRIAL USE

A hepatocyte growth factor has an activity to support the growth of immature myeloblast derived from mice. Furthermore, since the factor supports the growth of hemopoietic stem cells in an evaluation system using human bone marrow cells and murine bone marrow cells, it can be used as a stem cell augmenting agent having said hepatocyte growth factor as an active component for treatment of bone marrow suppression (for example, after the use of anti-cancer agents or after bone marrow transplantation), for treatment of bone marrow malfunctions (for example, hypoplastic anemia), or for in vitro growth of peripheral blood stem cells and bone marrow stem cells. Furthermore, since the hemopoietic stem cell augmenting agent in the present invention eventually promotes the growth of not only various blood cells but also osteoclasts which are progeny of hemopoietic stem cells, it is appropriately used as a therapeutic agent for osteoporosis or the like.

# SEQUENCE LISTING

5 10 15	SEQ ID NO: 1 SEQUENCE LENGTH: 16 SEQUENCE TYPE: amino acid TOPOLOGY: linear MOLECULE TYPE: peptide SEQUENCE DESCRIPTION:
20	Val'Val Asn Gly Ile Pro Thr Xaa Thr Asn Ile Gly 1 5 10
25	Xaa Met Val Lys 15
30	
35	
40	
45	
50	

SEQ ID NO: 2 SEQUENCE LENGTH: 2172 nucleic acid SEQUENCE TYPE: STRANDEDNESS: double TOPOLOGY: linear Genomic RNA cDNA to mRNA MOLECULE TYPE: ORIGINAL SOURCE: human normal fibroblast 10 SEQUENCE DESCRIPTION: 15 20 ATG TOG GTG ACC AAA CTC CTG CCA GCC CTG CTG CTG CAG CAT GTC CTC 48 Met Trp Val Thr Lys Leu Leu Pro Ala Leu Leu Cln His Val Leu 25 5 10 15 30 CTG CAT CTC CTG CTC CCC ATC GCC ATC CCC TAT GCA GAG GGA CAA 96 Leu His Leu Leu Leu Leu Pro Ile Ala Ile Pro Tyr Ala Glu Gly Gln 20 25 30 35 ACG AAA AGA AGA AAT ACA ATT CAT GAA TTC AAA AAA TCA GCA AAG ACT 144 Arg Lys Arg Arg Asn Thr Ile His Glu Phe Lys Lys Ser Ala Lys Thr 40 35 40 45 45 ACC CTA ATC AAA ATA GAT CCA GCA CTG AAG ATA AAA ACC AAA AAA GTG 192

50 55 60

Thr Leu Ile Lys Ile Asp Pro Ala Leu Lys Ile Lys Thr Lys Lys Val

	AAT	ACT	GCA	GAC	CAA	TCT	GCT	AAT	AGA	TGT	ACT	AGG	ААТ	AAA	CCA	CTT	240
	Asn	Thr	Ala	Asp	Gln	Cys	Ala	Asn	Arg	Cys	Thr	Arg	Asn	Lys	Gly	Leu	
5	65					70					75					80	
10	CCA	TTC	ACT	TCC	AAG	GCT	TTT	GTT	TTT	GAT	AAA	GCA	AGA	AAA	CAA	TGC	288
	Pro	Phe	Thr	Cys	Lys	Ala	Phe	Val	Phe	Asp	Lys	Ala	Arg	Lys	Gln	Cys	
					85			•		90					95		
15																	
	CTC	TCC	ŢŢĊ	$\alpha\alpha$	TTC	AAT	AGC	ATG	TCA	AGT	CGA	CTC	AAA	AAA	GAA	TTT	336
	Leu	Trp	Phe	Pro	Phe	Asn	Ser	Met	Ser	Ser	Gly	Val	Lys	Lys	Glu	Phe	
20				100				1	105				1	10			
25	CCC	CAT	GAA	TTT	GAC	CTC	TAT	GAA	AAC	AAA	GAC	TAC	ATT	AGA	AAC	TCC	384
	Gly	His	Glu	Phe	Asp	Leu	Tyr	Glu	Asn	Lys	Asp	Tyr	Ile	Arg	Asn	Cys	
			115				•	120				12	25				
30															-		
30	ATC	АТТ	GGT	AAA	GGA	ccc	AGC	TAC	AAG	GGA	ACA	GTA	TCT	ATC	- ACT	AAG	432
30			GGT Gly														432
30						Arg						Val					432
		Ile				Arg	Ser				Thr	Val					432
35	Ile	Ile 130		Lys	Gly	Arg	Ser 135	Tyr	Lys	Gly	Thr	Val 10	Ser	Ile	Thr	Lys	432 480
	Ile AGT	Ile 130 CGC	Gly	Lys AAA	Gly TGT	Arg CAG	Ser 135 CCC	Tyr TCG	Lys AGT	Gly	Thr 11 ATG	Val IO ATA	Ser CCA	Ile	Thr	Lys	
35	Ile AGT	Ile 130 CGC	Gly ATC	Lys AAA	Gly TGT Cys	Arg CAG	Ser 135 CCC	Tyr TCG	Lys AGT	Gly TCC Ser	Thr 11 ATG	Val IO ATA	Ser CCA	Ile	Thr	Lys CAC His	
35	Ile AGT Ser	Ile 130 CGC	Gly ATC	Lys AAA	Gly TGT Cys	Arg CAG Gln	Ser 135 CCC	Tyr TCG	Lys AGT	Gly TCC Ser	Thr 11 ATG Met	Val IO ATA	Ser CCA	Ile	Thr GAA Glu	Lys CAC His	
35 40	AGT Ser 145	Ile 130 GGC Gly	Gly ATC	AAA Lys	Gly TCT Cys	Arg CAG Gln 150	Ser 135 CCC Pro	Tyr TGG Trp	Lys AGT Ser	TCC Ser	Thr 11 ATG Met	Val 40 ATA Ile	Ser CCA Pro	Ile CAC His	Thr GAA Glu 160	CAC His	
35 40	AGT Ser 145	Ile 130 GGC Gly	Gly ATC Ile	AAA Lys CCT	Gly TGT Cys	CAG Gln 150	Ser 135 CCC Pro	Tyr TGG Trp CAG	AGT Ser	TCC Ser 19	Thr  ATG  Met  TAC	Val IO ATA Ile	Ser CCA Pro	CAC His	GAA Glu 160 CCT	CAC His	480
35 40	AGT Ser 145	Ile 130 GGC Gly	Gly ATC Ile	AAA Lys CCT	Gly TGT Cys	CAG Gln 150	Ser 135 CCC Pro	Tyr TGG Trp CAG	Lys AGT Ser GAA	TCC Ser 19	Thr  ATG  Met  TAC	Val IO ATA Ile	Ser CCA Pro	CAC His	GAA Glu 160 CCT	CAC His	480

	GGG	GAA	GAA	CCC	GGA	$\alpha$	TCC	TGT	TTC	ACA	ACC	AAT	CCA	GAG	GTA	ccc	576
5	Gly	Glu	Glu	Gly	Gly	Pro	Trp	Cys	Phe	Thr	Ser	Asn	Pro	Glu	Val	Arg	
				180				1	185				19	90			
10	TAC	GAA	GTC	TCT	GAC	ATT	CCT	CAG	TCT	TCA	GAA	CTT	GAA	TCC	ATG	ACC	624
	Tyr	Glu	Val	Cys	Asp	Ile	Pro	Gln	Cys	Ser	Glu	Val	Glu	Cys	Met	Thr	
15			195				2	200				20	05				
15																	
					AGT												672
20	Cys		Gly	Glu	Ser		_	Gly	Leu	Met	-		Thr	Glu	Ser	Gly	
		210				4	215				22	20					
	NCC.	ىتىدى	av•rı	CAC	ccc	ጥጥ	CAT	CAT	CAC	404	CCA	CAC	~~~	CAC		aviv-i	7720
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	TTG	CCT	GAA	AGA	TAT	ccc	GAC	AAG	CCC	TTT	GAT	GAT	AAT	TAT	TCC	ccc	768
					TAT Tyr												768
35					TAT Tyr 245				Gly						Cys		768
35					Tyr				Gly	Phe				Tyr	Cys		768
	Leu	Pro	Glu	Arg	Tyr	Pro	Asp	Lys	Gly 2	Phe 250	Asp	Asp	Asn	Tyr 25	Cys 55	Arg	768 816
35	Leu AAT	Pro CCC	Glu GAT	Arg CGC	Tyr 245	Pro CCG	Asp AGG	Lys	Gly Z	Phe 250 TCC	Asp	Asp ACT	Asn	Tyr 25 GAC	Cys 55 CCT	Arg CAC	
	Leu AAT	Pro CCC	Glu GAT	Arg CGC	Tyr 245 CAG	Pro CCG	Asp AGG	Lys CCA Pro	Gly Z	Phe 250 TCC	Asp	Asp ACT	Asn CTT Leu	Tyr 25 GAC	Cys 55 CCT	Arg CAC	
	Leu AAT	Pro CCC	Glu GAT	Arg GGC Gly	Tyr 245 CAG	Pro CCG	Asp AGG	Lys CCA Pro	Gly TCG Trp	Phe 250 TCC	Asp	Asp ACT	Asn CTT Leu	Tyr 25 GAC Asp	Cys 55 CCT	Arg CAC	
40	Leu AAT Asn	Pro CCC Pro	Glu GAT Asp	Arg GGC Gly 260	Tyr 245 CAG	Pro CCG Pro	Asp AGG Arg	CCA Pro	Gly TCG Trp 265	Phe 250 TCC Cys	Asp TAT Tyr	Asp ACT Thr	Asn CTT Leu 2	Tyr 25 GAC Asp 70	Cys 55 CCT Pro	Arg CAC His	
40 45	AAT Asn	Pro CCC Pro	Glu GAT Asp	Arg GGC Gly 260 GAG	Tyr 245 CAG Gln	Pro CCG Pro	Asp AGG Arg GCA	Lys CCA Pro	Gly TCG Trp 265	Phe 250 TGC Cys	Asp TAT Tyr	ASP ACT Thr	Asn CTT Leu 2	Tyr 25 GAC Asp 70	Cys 55 CCT Pro	Arg CAC His	816
40	AAT Asn	Pro CCC Pro	Glu GAT Asp TCG Trp	Arg GGC Gly 260 GAG	Tyr 245 CAG Gln TAC	Pro CCG Pro	Asp AGG Arg	CCA Pro	Gly TCG Trp 265	Phe 250 TGC Cys	Asp TAT Tyr	ASP ACT Thr	Asn CTT Leu 2	Tyr 25 GAC Asp 70	Cys 55 CCT Pro	Arg CAC His	816

	AAT	GAC	ACT	GAT	GTT	CCT	TTG	GAA	ACA	ACT	GAA	TCC	ATC	CAA	CCT	CAA	912
	Asn	Asp	Thr	Asp	Val	Pro	Leu	Glu	Thr	Thr	Glu	Cys	Ile	Gln	Gly	Gln	
5		290				2	295				30	00					
10	GGA	GAA	CCC	TAC	AGG	CCC	ACT	GTC	AAT	ACC	ATT	TCC	AAT	CGA	ATT	CCA	960
	Gly	Glu	Gly	Tyr	Arg	Gly	Thr	Val	Asn	Thr	Ile	Trp	Asn	Gly	Ile	Pro	
	305				3	310				3	15				320	)	
15																	
	TGT	CAG	CGT	TCG	GAT	TCT	CAG	TAT	CCT	CAC	GAG	CAT	GAC	ATG	ACT	CCT	1008
	Cys	Gln	Arg	Trp	Asp	Ser	Gln	Tyr	Pro	His	Glu	His	Asp	Met	Thr	Pro	
20					325				3	330				33	35		
25	GAA	AAT	TTC	AAG	TCC	AAG	GAC	CTA	CGA	GAA	AAT	TAC	TCC	CGA	AAT	CCA	1056
20	Glu	Asn	Phe	Lys	Cys	Lys	Asp	Leu	Arg	Clu	Asn	Tyr	Cys	Arg	Asn	Pro	
				340				3	345				35	50			
30															-		
	GAT	GGG	TCT	GAA	TCA	$\alpha$	TGG	TCT	TTT	ACC	ACT	GAT	CCA	AAC	ATC	CGA	1104
	Asp	Gly	Ser	Glu	Ser	Pro	Trp	Cys	Phe	Thr	Thr	Asp	Pro	Asn	Ile	Arg	
35			355				3	360				36	55				
10	GIT	GGC	TAC	TGC	TCC	CAA	ATT	CCA	AAC	TCT	GAT	ATG	TCA	CAT	GGA	CAA	1152
40	Val	Gly	Tyr	Cys	Ser	Gln	Ile	Pro	Asn	Cys	Asp	Met	Ser	His	Gly	Gln	
		370				3	375				38	30					
45																	
	GAT	TGT	TAT	CCT	CCC	AAT	GGC	AAA	AAT	TAT	ATG	GGC	AAC	TTA	TCC	CAA	1200
	Asp	Cys	Tyr	Arg	Gly	Asn	Gly	Lys	Asn	Tyr	Met	Gly	Asn	Leu	Ser	Gln	
50	385				3	390				39	95				400	כ	

	ACA	AGA	TCT	GGA	CTA	ACG	TGT	TCA	ATG	TGG	GAC	AAG	AAC	ATG	GAA	GAC	1248
5	Thr	Arg	Ser	Gly	Leu	Thr	Cys	Ser	Met	Trp	Asp	Lys	Asn	Met	Glu	Asp	
3					405				1	110				41	15		
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		Tyr															
20			435				ì	140				41	15	-	•	•	
25	ACG	GGA	AAT	CCA	CTC	ATT	CCT	TCC	GAT	TAT	TCC	CCT	ATT	TCT	CCT	TGT	1392
	Thr	Gly	Asn	Pro	Leu	Ile	Pro	Trp	Asp	Tyr	Cys	Pro	Ile	Ser	Arg	Cys	
		450				L	155				46	50					
30															-		
	GAA	GGT	GAT	ACC	ACA	CCT	ACA	ATA	CTC	AAT	TTA	GAC	CAT	$\alpha$	GTA	ATA	1440
	Glu	Gly	Asp	Thr	Thr	Pro	Thr	Ile	Val	Asn	Leu	Asp	His	Pro	Val	Ile	
35	465				1	170				47	75				480	)	
40	TCT	TGT	CCC	AAA	ACG	AAA	CAA	CTG	CGA	GTT	GTA	AAT	GGG	ATT	CCA	ACA	1488
	Ser	Cys	Ala	Lys	Thr	Lys	Gln	Leu	Arg	Val	Val	Asn	Gly	Ile	Pro	Thr	
					485				1	190				49	95		
45																	
	CGA	ACA	AAC	GTA	GGA	TCG	ATG	GTT	AGT	TTG	AGA	TAC	AGA	AAT	AAA	CAT	1536
	Arg	Thr	Asn	Val	Gly	Trp	Met	Val	Ser	Leu	Arg	Tyr	Arg	Asn	Lys	His	
50				5 <b>0</b> 0				9	505				5	10			

	ATC	TCC	CGA	GGA	TCA	MG	ATA	AAG	GAG	ACT	TCG	CTT	CTT	ACT	GCA	CGA	1584
	Ile	Cys	Gly	Gly	Ser	Leu	Ile	Lys	Glu	Ser	Trp	Val	Leu	Thr	Ala	Arg	
5			515				. 5	520				52	25				
10	CAG	TGT	TTC	CCT	TCT	CGA	GAC	TTG	AAA	GAT	TAT	GAA	CCT	TGG	CTT	GGA	1632
	Gln	Cys	Phe	Pro	Ser	Arg	Asp	Leu	Lys	Asp	Tyr	Glu	Ala	Trp	Leu	Gly	
		530					535				51	10					
15																	
	ATT	CAT	GAT	GTC	CAT	GGA	AGA	CGA	GAT	GAG	AAA	TCC	AAA	CAG	GTT	CTC	1680
	Ile	His	Asp	Val	His	Gly	Arg	Gly	Asp	Glu	Lys	Cys	Lys	Gln	Val	Leu	
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25	AAT	GTT	TCC	CAG	CTG	GTA	TAT	ccc	CCT	GAA	CGA	TCA	GAT	CTG	CTT	TTA	1728
	Asn	Val	Ser	Gln	Leu	Val	Tyr	Gly	Pro	Glu	Gly	Ser	Asp	Leu	Val	Leu	
					565				ģ	570				5'	75		
30																	
	ATG	AAG	CTT	GCC	AGG	CCT	GCT	GTC	CTG	GAT	GAT	TTT	GTT	AGT	ACG	ATT	1776
	Met	Lys	Leu	Ala	Arg	Pro	Ala	Val	Leu	Asp	Asp	Phe	Val	Ser	Thr	Ile	
35	Met	Lys	Leu	Ala 580	Arg	Pro	Ala		Leu 585	Asp	Asp	Phe		Ser 90	Thr	Ile	
35	Met	Lys	Leu		Arg	Pro	Ala			Asp	Asp	Phe			Thr	Ile	
35								ŗ.	585				59	90			1824
	GAT	TTA	CCT	580 AAT	TAT	GGA	TGC	ACA	585 ATT	CCT	GAA	DAA	59 ACC	90 AGT	TCC		1824
	GAT	TTA	CCT	580 AAT Asn	TAT	GGA	TGC Cys	ACA	585 ATT	CCT	GAA	AAG Lys	59 ACC	90 AGT	TCC	AGT	1824
	GAT	TTA	CCT Pro	580 AAT Asn	TAT	GGA	TGC Cys	ACA Thr	585 ATT	CCT	GAA	AAG Lys	59 ACC Thr	90 AGT	TCC	AGT	1824
40	GAT Asp	TTA Leu	CCT Pro 595	580 AAT Asn	TAT Tyr	GGA Gly	TGC Cys	ACA Thr 600	ATT	CCT Pro	GAA Glu	AAG Lys	59 ACC Thr 05	90 AGT Ser	TCC Cys	AGT	1824 1872
40	GAT Asp	TTA Leu	CCT Pro 595	580 AAT Asn	TAT Tyr	GGA Gly TAC	TGC Cys	ACA Thr 600 GGA	ATT Ile	CCT Pro	GAA Glu AAC	AAG Lys 6	ACC Thr 05	AGT Ser	TCC Cys	ACT Ser	
40	GAT Asp	TTA Leu	CCT Pro 595	580 AAT Asn	TAT Tyr	GGA Gly TAC Tyr	TGC Cys	ACA Thr 600 GGA	ATT Ile	CCT Pro	GAA Glu AAC Asn	AAG Lys 6	ACC Thr 05	AGT Ser	TCC Cys	ACT Ser	

	CGA	GTG	CCA	CAT	CTC	TAT	ATA	ATG	CGA	TAA	GAG	AAA	TCC	AGC	CAG	CAT	1920
	Arg	Val	Ala	His	Leu	Tyr	Ile	Met	Gly	Asn	Glu	Lys	Cys	Ser	Gln	His	
5	625				$\epsilon$	30				63	35				640	)	
10	CAT	CGA	GGG	AAG	GTG	ACT	CTG	AAT	GAG	TCT	GAA	ATA	TGT	CCT	CCC	CCT	1968
10	His	Arg	Gly	Lys	Val	Thr,	Leu	Asn	Glu	Ser	Glu	Ile	Cys	Ala	Gly	Ala	
					645				6	550				6	55		
15																	
	GAA	AAG	ATT	GGA	TCA	GGA	CCA	TGT	GAG	CCC	GAT	TAT	CCT	CCC	CCA	CTT	2016
	Glu	Lys	Ile	Gly	Ser	Gly	Pro	Cys	Glu	Gly	Asp	Tyr	Gly	Gly	Pro	Leu	672
20				660				(	665				6'	70			
	GTT	TGT	GAG	CAA	CAT	AAA	ATG	AGA	ATG	GTT	CTT	CCT	CTC	ATT	GTT	CCT	2064
25	Val	Cys	Glu	Gln	His	Lys	Met	Arg	Met	Val	Leu	Gly	Val	Ile	Val	Pro	
			675				6	580				68	35				
30															-		
	CCT	CGT	GGA	TGT	CCC	ATT	CCA	AAT	CGT	CCT	CCT	ATT	TTT	GTC	CGA	GTA	2112
	Gly	Arg	Gly	Cys	Ala	Ile	Pro	Asn	Arg	Pro	Gly	Ile	Phe	Val	Arg	Val	
35		690				(	595				70	00					
40	GCA	TAT	TAT	GCA	AAA	TCC	ATA	CAC	AAA	ATT	ATT	TTA	ACA	TAT	AAG	GTA	2160
40	Ala	Tyr	Tyr	Ala	Lys	Trp	Ile	His	Lys	Ile	Ile	Leu	Thr	Tyr	Lys	Val	
	705	;				710				7	15				72	0	
45																	
	CCA	CAG	TCA	TAG													2172
	Pro	Gln	Ser	***													
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## Claims

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1. A hemopoietic stem cell augmenting agent comprising at least one hepatocyte growth factor as an active component.

- 2. A hemopoietic stem cell augmenting agent according to claim 1 wherein interleukin 3 and/or interleukin 7 are contained additionally as an active component(s).
- 3. A hemopoietic stem cell augmenting agent according to claim 1 which is used as a therapeutic agent for bone marrow suppression.
- 4. A hemopoietic stem cell augmenting agent according to claim 1 which is used as a therapeutic agent for bone marrow malfunctions.
- 5. A hemopoietic stem cell augmenting agent according to claim 1 wherein the hepatocyte growth factor is a protein which has a molecular weight of about 60,000 and has the following N-terminal amino acid sequence:

Val Val Asn Gly lle Pro Thr Xaa Thr Xaa Thr Asn lle Gly Xaa Met Val Lys (wherein Xaa is a conventional amino acid residue.)

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- 6. A hemopoietic stem cell augmenting agent according to claim 5 wherein the hepatocyte growth factor is a protein which has the amino acid composition of Table 1.
- 7. A hemopoietic stem cell augmenting agent according to claim 1 wherein the hepatocyte growth factor is a recombinant-type hepatocyte growth factor derived from human hepatic cells, a recombinant-type hepatocyte growth factor derived from fibroblasts or a hepatocyte growth factor derived from the human placenta or an equivalent substance in terms of effectiveness.
- 8. A hemopoietic stem cell augmenting agent according to claim 1 wherein the hepatocyte growth factor is obtained by purification using chromatography in which a gel carrier is bound with zinc via chelate.
  - 9. A hemopoietic stem cell augmenting agent according to claim 8 characterized in that a carrier bound additionally with heparin is used as the carrier.
- 10. A hemopoietic stem cell augmenting agent according to claim 1 wherein the hepatocyte growth factor is a protein having the amino acid sequence shown in SEQ ID NO:2 of Sequence Listing, or an equivalent substance in terms of effectiveness.
- 11. A process for the augmentation of hemopoietic stem cells characterized in that immature bone marrow cells are differentiated and proliferated in the presence of at least one hepatocyte growth factor.
  - **12.** A process for the augmentation of hemopoietic stem cells according to claim 11 characterized in that pluripotent hemopoietic stem cells are differentiated and proliferated.
- 40 **13.** A process for the augmentation of hemopoietic stem cells according to claim 11 characterized in that human bone marrow cells obtained from peripheral blood stem cells or bone marrow fluid.
  - 14. A protein which is a type of hepatocyte growth factor and has hemopoietic stem cell augmenting activity.

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- **15.** A protein according to claim 14 wherein the protein has a molecular weight of about 60,000 and has the following N-terminal amino acid sequence:
  - Val Val Asn Gly lle Pro Thr Xaa Thr Xaa Thr Asn lle Gly Xaa Met Val Lys (in which Xaa is a conventional amino acid.)

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- 16. A protein according to claim 5 wherein the protein has the amino acid composition of Table 1.
- 17. A protein according to claim 14 wherein the protein is obtained by purification through chromatography using a carrier bound with a zinc-chelate group.

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18. A protein according to claim 17 characterized in that a carrier bound additionally with heparin is used as the carrier.

- **19.** A protein according to claim **14** wherein the protein is an immature bone marrow cell growth factor and is a type of hepatocyte growth factor.
- 20. A protein according to claim 14 wherein the protein has a pluripotent hemopoietic stem cells augmenting activity and is a type of hepatocyte growth factor and a recombinant human hepatocyte growth factor.

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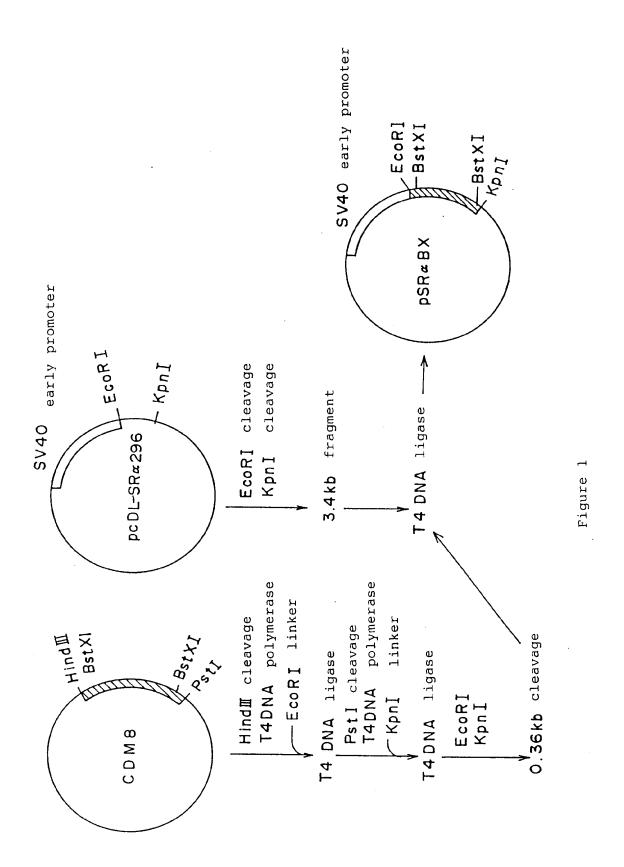
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- 21. A protein according to claim 14 wherein the protein has the amino acid sequence shown in SEQ ID NO:2 of Sequence Listing, or an equivalent substance in terms of effectiveness.
- 22. A protein according to claim 21 wherein said recombinant human hepatocyte growth factor is the one expressed in CHO cells derived from the Chinese hamster or COS cells derived from the monkey.
- 23. A DNA characterized in that it contains a nucleotide sequence which encodes a protein which is a type of hepatocyte growth factor and has a hemopoietic stem cell augmenting activity or an equivalent substance in terms of effectiveness.
  - 24. A DNA according to claim 23 wherein said amino acid sequence has the amino acid sequence shown in SEQ ID NO:2 of Sequence Listing, or an equivalent substance in terms of effectiveness.
  - 25. A recombinant expression vector in which a nucleotide sequence encoding an amino acid sequence of a protein which is a type of hepatocyte growth factor and has a hemopoietic stem cell augmenting activity or an equivalent substance thereof in terms of effectiveness is incorporated in such a manner that it can be expressed.
  - 26. A transformant which is obtained by transforming a host cell with a recombinant expression vector in which nucleotide sequence encoding an amino acid sequence of a protein which is a type of hepatocyte growth factor and has a hemopoietic stem cells augmenting activity or an equivalent substance thereof in terms of effectiveness, is incorporated in such a manner that it can be expressed.
  - 27. A transformant according to claim 26 wherein said transformant is <u>E. coli</u> or cells derived from mammalians.
- 28. A process for the preparation of a recombinant human hepatocyte growth factor protein, which comprises transforming a host cell with a recombinant expression vector in which a nucleotide sequence coding for an amino acid sequence of a hepatocyte growth factor having a hemopoietic stem cell augmenting activity or an equivalent substance thereof in terms of effectiveness is incorporated expressibly, to produce a transformant, culturing the resultant transformant under such conditions that said protein or equivalent substance can be expressed in a nutrient medium, to produce said recombinant human hepatocyte growth factor protein and collecting and isolating the produced recombinant human hepatocyte growth factor protein from said culture.



# Figure 2 (1)

			ACC Thr													48
1			5	5				10	)				5			
			CTC Leu					Ala					Glu			96
			20					25					30			
			AGA Arg													144
ACC	СТА	ATC	AAA	ATA	GAT	<b>∝</b> A	CCA	CTG	AAG	АТА	AAA	ACC	AAA	AAA	GTG	192
Thr	Leu 50	Ile	Lys	Ile	Asp	Pro 55	Ala	Leu	Lys	Ile	Lys 60	Thr	Lys	Lys	Val	
AAT	ACT	CCA.	GAC	CAA	TGT	cct	AAT	AGA	TGT	ACT	ACG	AAT	AAA	CGA	CTT	240
Asn 65	Thr	Λla	Asp	Gln	Cys 70	Ala	Asn	Arg	Cys	Thr 75	Arg	Asn	Lys	Gly	Leu 80	
CCA.	TTC	ACT	TCC	ΛAG	CCT	TTT	CIT	TTT	GAT	AAA	CCA	AGA	AAA	CAĀ	TCC	288
Pro	Phe	Thr	Cys	Lys 85	Ala	Phe	Val	Phe	Asp 90	Lys	Ala	Arg	Lys	G1n 95	Cys	
CIC	TCC	TTC	$\infty$	TTC	AAT	ACC	ATG	TCA	AGT	CCA	GTG	AAA	AAA	GAA	TTT	336
Leu	Trp	Phe	Pro 100	Phe	Asn	Ser		Ser 105	Ser	Gly	Val	· ·	Lys 10	Glu	Phe	
$\infty$	CAT	GAA	TTT	GAC	CTC	TAT	GAA	AAC	AAA	GAC	TAC	ATT	AGA	AAC	TOC	384
Gly	His	Glu 115		Asp	Leu	_	G1u 120	Asn	Lys	Asp	_	Ile 25	Arg	, Asn	Cys	
ATC	ATT	œī	AAA	CGA	$\alpha$	ACC	TAC	AAG	CGA	ACA	GTA	TCT	ATC	ACT	AAG	432
Ile	Ile	-	Lys	Gly	_	Ser	Tyr	Lys	Gly		Val	Ser	Ile	Thr	Lys	

# Figure 2 (2)

AGT	000	ATC	: AAA	TGT	CAC	$\infty$	TOC	ACT	TCC	ATG	АТА	CCA	CAC	GAA	CAC	480
Ser	Gly	Ile	Lys	Cys	Gln	Pro	Trp	Ser	Ser	Met	Ile	Pro	His	Glu	His	
145					150	-			1	55				16	0	
															CGA	528
Ser	Tyr	Arg	Gly	Lys	Asp	Leu	Gln	Glu	Asn	Tyr	Cys	Arg	Asr	i Pro	Arg	
				165	5				170				1	75		
															$\alpha$	576
Gly	Glu	Glu			Pro	Trp			Thr	Ser	· Asn	Pro	Glu	ı Val	Arg	
			180	l				185				1	90			
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# Figure 2 (3)

AAT GAC ACT GAT GTT CCT TTG GAA ACA ACT GAA TCC ATC CAA CGT CAA

Asn	Asp	Thr	Asp	Val	Pro	Leu	Glu	Thr	Thr	Glu	Cys	Ile	Gln	Gly	Gln	
	290				2	95				30	0					
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					Lys											
			340				,	345				3	50			
GAT	œ	TCT	GAA	TCA	$\infty$	TCG	TGT	TTT	ACC	ACT	GAT	CCA	AAC	ATC	CGA	1104
Asp	Gly	Ser	Glu	Ser	Pro	Trp	Cys	Phe	Thr	Thr	Asp	Pro	Asn	Ile	Arg	
		355					360				3	65				
														_		
															CAA	1152
Val	Gly	Tyr	Cys	Ser	Gln	Ile	Pro	Asn	Cys	Asp	Met	Ser	His	Gly	Gln	
	370					375				3	80					
													. mna	mv-v		1200
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-	_	Tyr	Are	GLy		Gly	Lys	asn.			: GI?	/ Asr	Leu		Gln	
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															u Asp	
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TT	CAC		CAT	OTA 7	c mo	c TOX	G GA	A OCA	\ GAT	r oc <i>i</i>	A ACT	r aac	CIO	G AA'	r gag	1296
															n Glu	
			420			_		425					430			

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# Figure 2 (4)

ΑΛΤ	TAC	TCC	CCA	TAA	CCA	GAT	GAT	GAT	CCT	CAT	CCA	$\infty$	TCC	TCC	TAC	1344
Asn	Tyr	Cys	Arg	Asn	Pro	Asp	Asp	Asp	Ala	His	Gly	Pro	Trp	Cys	Tyr	
		435				Ц	40				41	15				
												ATT				1392
Thr	_	Asn	Pro	Leu	_		Trp	Asp	Tyr			Ile	Ser	arg	Cys	
	450					155				40	60					
GAA	CCT	GAT	ΛŒ	ACA	CCT	ACA	ATA	GIC	AAT	TTA	GAC	CAT	$\infty$	GTA	ATA	1440
												His				
465	_				470					75				48		
TCT	TCT	$\infty$	AAA	ΛCG	AAA	CAA	CTG	CGA	GIT	GTA	AAT	$\alpha$	ATT	CCA.	ACA	1488
Ser	Cys	Ala	Lys	Thr	Lys	Gln	Leu	Arg	Val	Val	Asn	Gly	Ile	Pro	Thr	
				485					490				14	95		
															0 A //	1506
															CAT	1536
Arg	Thr	Asn		Gly	Trp	Met			Leu	Arg	Tyr		_	Lys	His	
			500					505				כ	10			
ΔΤΥ	TYY	CCA	CCA.	TCA	TTY	ΔΤΔ	AAG	GAG	AGT	1755	GTT	CTT	ACT	CCA	CGA	1584
															Arg	
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CAG	TGT	TTC	CT	TCT	· CGA	GAC	TTC	AAA	GAT	TAT	GAA	CCT	TOO	CM	CGA	1632
Gln	Cys	Phe	Pro	Ser	Are	Asp	Leu	ı Lys	Asp	Туг	Glu	ı Ala	Trp	Leu	Gly	
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#### EP 0 550 769 A1

## Figure 2 (5)

				ACC												1776
Met	Lys	Leu		Arg	Pro	Ala		_	Asp	Asp	Phe	Val	Ser	Thr	Ile	
			580					585				59	90			
GAT	ТТА	CCT	АЛТ	TAT	CGA	TCC	ACA	ልሞተ	CCT	CAA	AAC:	۸۲۲	٨	יציזו	۸۲۳	1020
				Tyr												1824
		595			·		600					05		-,-	20.	
				ccc												1872
Val			Trp	Gly			Gly	Leu	Ile	Asn	Tyr	Asp	Gly	Pro	Leu	
	610	:			(	615				62	20					
CGA	GIG	CCA	САТ	CTC	ТАТ	ΔΤΔ	ΔΤΥ	CCA	ΔΔΤ	CAC	ΔΛΛ	יצאו	۸۵۲	CAG	CAT	1920
				Leu												1920
625					530			J		35	•	3		640		
CAT	CCA	$\infty$	AAG	GTG	ACT	CTG	AAT	GAG	TCT	GAA	ATA	TCT	CCT	occ	CCT	1968
His	Arg	Gly	Lys	Val	Thr	Leu	Asn	Glu	Ser	Glu	Ile	Cys	Ala	Gly	Ala	
				645				6	550				65	55		
GAA	AAG	ТТА	GGA		CCGA.	CCA.	TGT			GAT	ТАТ	aar		•	لملت	2016
				TCA				GAG	œ				œc	CCA		2016 672
		Ile					Cys	GAG	œ				cc Gly	CCA		
Glu	Lys	Ile	Gly 660	TCA Ser	Gly	Pro	Cys 6	GAG Glu 665	CCC Gly	Asp	Tyr	Gly 67	GCC Gly 70	CCA Pro	Leu	
Glu	Lys TGT	Ile GAG	Gly 660 CAA	TCA Ser CAT	Gly AAA	Pro ATG	Cys 6 AGA	GAG Glu 565 ATG	CCC Gly GTT	Asp CTT	Tyr OGT	Gly 67 GIC	GCC Gly O	CCA Pro	Leu CCT	
Glu	Lys TGT	Ile GAG Glu	Gly 660 CAA	TCA Ser	Gly AAA	Pro ATG Met	Cys 6 AGA Arg	GAG Glu 565 ATG	CCC Gly GTT	Asp CTT	Tyr CCT Gly	Gly 67 GTC Val	GCC Gly O	CCA Pro	Leu CCT	672
Glu	Lys TGT	Ile GAG	Gly 660 CAA	TCA Ser CAT	Gly AAA	Pro ATG Met	Cys 6 AGA	GAG Glu 565 ATG	CCC Gly GTT	Asp CTT	Tyr OGT	Gly 67 GTC Val	GCC Gly O	CCA Pro	Leu CCT	672
Glu GTT Val	Lys TGT Cys	Ile GAG Glu 675	Gly 660 CAA Gln	TCA Ser CAT His	Gly AAA Lys	Pro ATG Met	Cys 6 AGA Arg 580	GAG Glu 565 ATG Met	CCC Gly CTT Val	Asp CTT Leu	Tyr CCT Gly 68	Gly 67 GIC Val 85	GCC Gly 'O ATT Ile	CCA Pro GTT Val	Leu CCT Pro	672 2064
Glu GTT Val	Lys TGT Cys	GAG Glu 675 GGA	Gly 660 CAA Gln TGT	TCA Ser CAT	Gly AAA Lys ATT	ATG Met CCA	Cys AGA Arg 580 AAT	GAG Glu 565 ATG Met	CCT	Asp CTT Leu	Tyr CCT Gly 68 ATT	Gly 67 GTC Val 35	GCC Gly 70 ATT Ile	CCA Pro	Leu CCT Pro	672
Glu GTT Val GGT Gly	Lys TGT Cys	GAG Glu 675 GGA	Gly 660 CAA Gln TGT	TCA Ser CAT His	Gly AAA Lys ATT Ile	ATG Met CCA	Cys AGA Arg 580 AAT	GAG Glu 565 ATG Met	CCT	Asp CTT Leu	CCT Cly 60 ATT Ile	Gly 67 GTC Val 35	GCC Gly 70 ATT Ile	CCA Pro	Leu CCT Pro	672 2064
Glu GTT Val GGT Gly	TGT Cys CGT Arg 690	GAG Glu 675 GGA Gly	Gly 660 CAA Gln TGT Cys	TCA Ser CAT His	AAAA Lys ATT Ile	ATG Met CCA Pro	Cys AGA Arg 580 AAT Asn	GAG Glu 665 ATG Met CGT Arg	CCT-Pro	Asp CTT Leu CCT Gly 70	CCT Cly 68 ATT Ile	Gly GTC Val 35 TTT Phe	CCC Gly 70 ATT Ile CTC Val	CCA Pro GTT Val	Leu CCT Pro CTA Val	672 2064
Glu GTT Val GGT Gly CCA	TGT Cys CGT Arg 690	GAG Glu 675 GGA Gly	Gly 660 CAA Gln TGT Cys	TCA Ser CAT His	AAAA Lys ATT Ile	ATG Met CCA Pro 695	Cys AGA Arg 580 AAT Asn	GAG Glu 665 ATG Met	CCT Pro	Asp CTT Leu CCT Gly 70	Tyr CCT Gly 68 ATT Ile	Gly 67 GTC Val 35 TTT Phe	CCCC Gly 'O ATT Ile GTC Val	CCA Pro GTT Val CGA Arg	CCT Pro CTA Val	672 2064
Glu GTT Val GGT Gly CCA	TGT Cys CGT Arg 690	GAG Glu 675 GGA Gly	Gly 660 CAA Gln TGT Cys	TCA Ser  CAT His  CCC Ala  AAA Lys	AAAA Lys ATT Ile	ATG Met CCA Pro 695	Cys AGA Arg 580 AAT Asn	GAG Glu 665 ATG Met	CCT Pro	Asp CTT Leu CGT Gly 70 ATT Ile	Tyr CCT Gly 68 ATT Ile	Gly 67 GTC Val 35 TTT Phe	CCCC Gly 'O ATT Ile GTC Val	CCA Pro GTT Val CGA Arg	CCT Pro CTA Val	672 2064 2112

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Figure 2 (6)

CCA CAG TCA TAG
Pro Gln Ser \*\*\*

2172

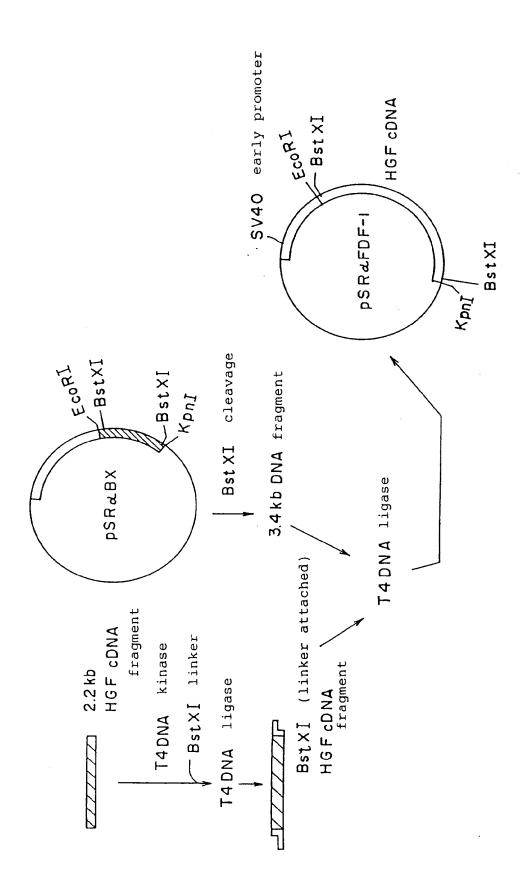


Figure 3

NFS60 Cell Growth
Stimulating Activity (U/ml)

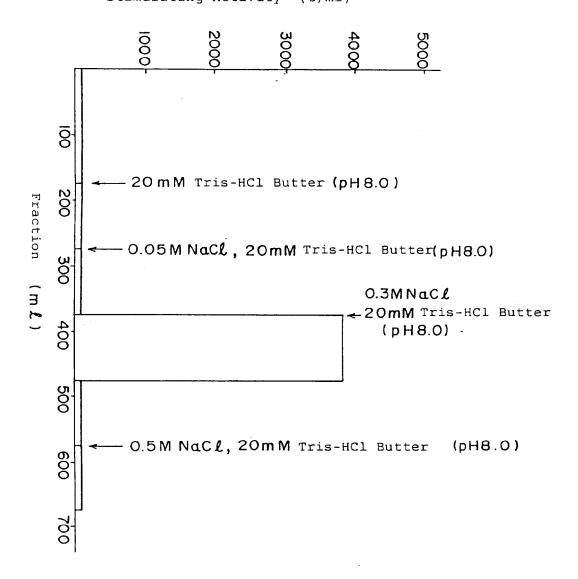


Figure 4

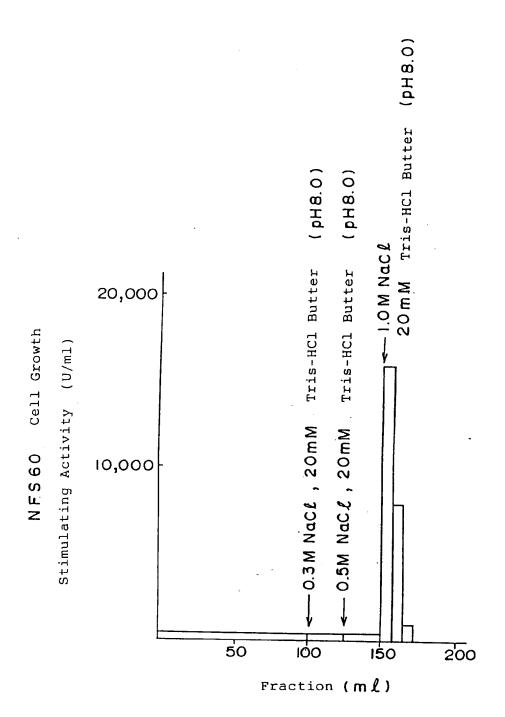


Figure 5

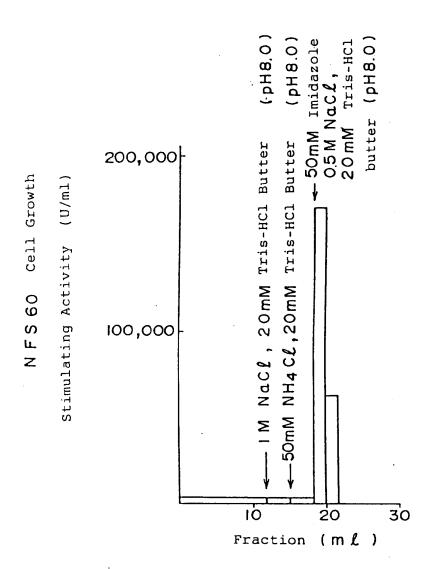


Figure 6

Molecular Marker Purified Recombinant HGF

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97 —\_\_

66 \_\_\_

45 —

31 —

21.5 \_\_\_

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Figure 7

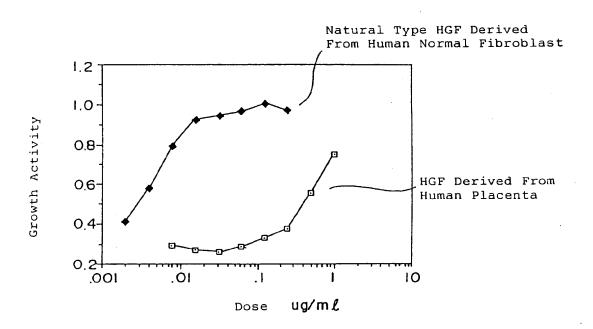
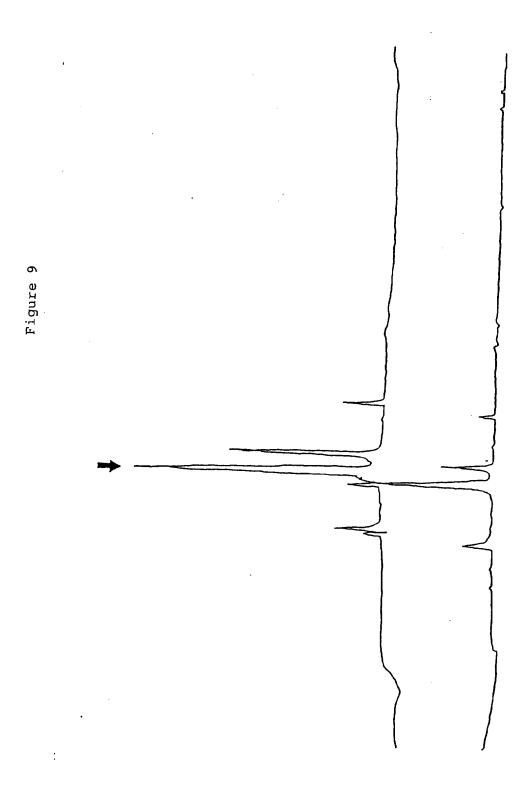


Figure 8



## INTERNATIONAL SEARCH REPORT

International Application No PCT/JP92/00949

			International Application No PCT	/JP92/00949					
I. CLASS	SIFICATIO	N OF SUBJECT MATTER (if several class	ification symbols apply, indicate all) 6						
		tional Patent Classification (IPC) or to both Na							
Int —	. Cl <sup>5</sup>	C07K15/04, C12N15/18	3, C12P21/02, A61K3	7/02					
II. FIELD	S SEARC	HED							
		Minimum Docume	ntation Searched 7						
Classificati	on System		Classification Symbols						
		C07K15/04, 15/06, C1	2N1E/12 1E/10						
IPO	2	15/27, C12P21/00, 21	12N13/12, 13/18,						
		1							
		Documentation Searched other to the Extent that such Documents	than Minimum Documentation s are Included in the Fields Searched s						
Bio	logica	al Abstracts Data Base	(BTOSTS)						
Cher	nical	Abstracts Data Base	(CA. REGISTRY)						
		The Control of Salar Salar	(City REGISTRI)						
III. DOCU	MENTS C	ONSIDERED TO BE RELEVANT							
Category •	Citat	ion of Document, 11 with indication, where app	propriate, of the relevant passages 12	Relevant to Claim No. 13					
A	Proc	ceedings of the Nation	al Academy of	1-28					
	Scie	ences of the USA, Vol.	84, No. 24,	1 20					
	(198	37), K. Ikebuchi et al	-•,						
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"A" docu	ment defin	ing the general state of the art which is not	"T" later document published after the priority date and not in conflict with	the application but cited to					
		e of particular relevance at but published on or after the international	understand the principle or theory "X" document of particular relevance; t						
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whic	h is cited t	n may throw doubts on priority claim(s) or o establish the publication date of another	"Y" document of particular relevance; the claimed invention cannot be considered to involve an invention standard the document						
		special reason (as specified) ing to an oral disclosure, use, exhibition or	be considered to involve an inventive step when the documer is combined with one or more other such documents, suc combination being obvious to a person skilled in the art						
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IV. CERTI									
		mpletion of the International Search	Date of Mailing of this International Sea	arch Report					
		, 1992 (19. 10. 92)	November 2, 1992 (02. 11. 92)						
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V. OBSERVATIONS WHERE CERTAIN CLAIMS WERE FOUND UNSEARCHABLE '	:
This international search report has not been established in respect of certain claims under Article 17(2) (a) to 1. Claim numbers . because they relate to subject matter not required to be searched by this	•
2. Claim numbers , because they relate to parts of the international application that do not correquirements to such an extent that no meaningful international search can be carried out, specific	nply with the prescribed cally:
3. Claim numbers , because they are dependent claims and are not drafted in accordance with sentences of PCT Rule 6.4(a).	th the second and third
VI OBSERVATIONS WHERE UNITY OF INVENTION IS LACKING 2	
This International Searching Authority found multiple inventions in this international application as follows:	ws:
1. As all required additional search fees were timely paid by the applicant, this international search rep claims of the international application	ort covers all searchable
2. As only some of the required additional search fees were timely paid by the applicant, this international those claims of the international application for which fees were paid, specifically claims:	search report covers only
3. No required additional search fees were timely paid by the applicant. Consequently, this international search fees were timely paid by the applicant. Consequently, this international search fees were timely paid by the applicant. Consequently, this international search fees were timely paid by the applicant. Consequently, this international search fees were timely paid by the applicant. Consequently, this international search fees were timely paid by the applicant. Consequently, this international search fees were timely paid by the applicant. Consequently, this international search fees were timely paid by the applicant.	irch report is restricted to
4. As all searchable claims could be searched without effort justifying an additional fee, the International Se invite payment of any additional fee.	arching Authority did not
Remark on Protest  The additional search fees were accompanied by applicant's protest.	·
No protest accompanied the payment of additional search fees.	